AA505075-derived oligonucleotide SEQ ID 4947.

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Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                     bronchodilating agent.
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, Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sandrasagra A,
L, Shahabuddin
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S Katz E,

Pabalan J,

Aguilar

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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, comprising allergies are derived from a gene encoding or regulating continuous expression of a target polypeptide associated with lung airway or lung captures, in separate containers, (b) the oligonucleotides, (c) device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, confit in the invention has antiallergic, antiinflammatory, antiasthmatic, confit ending a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The composition composition, allergies and/or bronchoconstriction are associated with a disease or condition such as pulmonary vasoconstriction, confit respiratory, and/or bronchoconstriction and/or lung confit respiration, emphysema, chronic obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary confit respiration, respiratory pulmonary path, cystic fibonesis, allergic rhinitis, pulmonary to the antial respiration, pulmonary disease, pulmonary confit respiration in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system confit and affects disease, the circular free adenosine into the system confit the confit of th SEQ ID NO 4947; 763pp; English. unwanted effects due to it

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

Matches Query Match Local 16; Similarity Conservative 1.1%; 0 Score 14.8; Pred. No. 4. Mismatches 4.4e+02; DB 1; Length Indels 0

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1520 AAAAAAAAAAAAGTAAAA 1537

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ABD25936 standard; DNA; 20

ABD25936;

29-JUL-2004 (first entry)

AA505075-derived oligonucleotide SEQ ID 4948

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; pulmonary transplantation rejection; ss; primer. cancer; bronchitis;

Homo sapiens.

WO200285309-A2.

31-OCT-2002.

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23-APR-2002; 2002WO-US013143

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC

Nyce JW, Miller S, Ś Li Y, Tang Sandrasagra A, L, Shahabuddin ŝ Katz 'n Pabalan ۲, Aguilar

'n

WPI; 2003-093058/08

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, target nucleic acids associated with lung airway or lung dysfunction, bronchodilating agent. targeted to

Claim 15; SEQ ID NO 4948; 763pp; English.

RESULT 752
ABD25936
ID ABD259
XX ABD259
XX ABD259
XX ABD259
XX ABD259
XX ABD259
XX AA5050
XX Human;
XX Feepir
XX Empir
XX Empir
XX Empir
XX Pulmon
XX Homo s
XX W Pulmon
XX Fee
PT Nis 1:
CC Compori
CC Surfac
CC Consoli
CC Greduci
CC CC comprising oligonucleotides, effective for alleviating composition of ligonucleotides, effective for alleviating controlled adenosine sensitivity, levels of adenosine (A) or (A) receptors, considered and sensitivity, levels of adenosine (A) or (A) receptors, considered and sensitivity, levels of adenosine (A) or (A) receptors, considered and sensitivity, levels of adenosine (A) or (A) receptors, considered and sensitivity, and considered with lung airway or lung considered considered with allevery (C) analysis, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, is a letter addrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The gulmonary obstruction, and/or bronchoconstriction and/or lung constitution, and/or bronchoconstriction are associated with a disease or condition such as pulmonary vasoconstriction, comprises only a pain, cystic fiboresis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary cancer.

The constraint of the constructive pulmonary disease, pulmonary cancer. transplantation rejection, pulmonary infections, bronce The reduced adenosine content of the anti-sense oligos This invention describes a novel composition (a) a first active agent,

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ABD32081/c
ID ABD320
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AC ABD320
XX ABD320
XX 29-JUL
XX Human
XX Human
XX respir
KW respir
KW respir
KW respir
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Best Local S
Matches 16
                   comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, insuppressive, immunosuppressive, and cyrostatic activity, is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nyce JW,
Miller S
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the oligonucleotides into pro
e.g., lung, brain, heart, kid
prevent any unwanted effects
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                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel composition (a) a
                                                                                                                                                                                                                                                                                                                                                                                                             Claim 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            bronchodilating agent.
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16; Conserv
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Tang
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                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 14292; 763pp; English.
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L, Shahabuddin
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Pred. No. 4
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. 4.4e+02;
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           for
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RESULT 754
ABD215/C
ID ABD215
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ABD216
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Human;
KW respir
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                                                                                                                                                                   Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targenucleic acids associated with lung airway or lung dysfunction
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                                                                                                                                           agent.
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L, Shahabuddin
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Claim

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SEQ ID NO

553;

763pp; English.

comprising

oligonucleotides,

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novel composition (a) a firs, effective for alleviating

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active agent,

This invention describes

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CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a CC beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered CC composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to CC reduce the amount of target polypeptide present in the lungs. The CC pulmonary obstruction, and/or bronchoconstriction and/or lung CC inflammation, allergies and/or surfactant hypoproduction are associated CC with a disease or condition such as pulmonary vasoconstriction, cepitatory CC distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary CC hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary CC transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the charget RNA serves to prevent the breakdown of the charget RNA serves to prevent the breakdown of the charget RNA serves to prevent the breakdown of the charget RNA serves to prevent the breakdown of the charget RNA serves to prevent the breakdown of the charget RNA serves to prevent the breakdown of the charget RNA serves to prevent the breakdown of the charget RNA serves to prevent the breakdown of the charget RNA serves to prevent the breakdown of the charget RNA serves to prevent the breakdown of the charget RNA serves to prevent the prevent
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Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                pulmonary transplantation rejection;
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ilarity 88.9%;
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L, Shahabuddin
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                                                                                            PHARM INC.
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Pred. No. 4.4e
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cc comprising oligonucleotides, effective for alleviating cc bronchoconstriction, respiratory tract inflammation, allergies and ceducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, converged to a mammal. The coligonucleotides are derived from a gene encoding or regulating cc expression of a target polypeptide associated with lung airway or lung cystuction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, consposition comprises oligo and is antiinflammatory, antiasthmatic, can algesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the mount of target polypeptide present in the lungs. The composition of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, allergies and/or bronchoconstriction and/or lung inflammation, allergies and/or bronchoconstriction and/or lung inflammation, allergies, asthma, impeded respiration, respiratory continues syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary tractory disease, pulmonary mifections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the coligonary brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                RESULT 75
ABD21765
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a novel composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 15; SEQ ID NO 4683;
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Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

sapiens

Human stanniocalcin-derived oligo SEQ

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ABD21765;

29-JUL-2004

(first entry)

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CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or composition comprises oligo and is administered to reduce the production composition, languaged to inflammation, altergies and/or bronchoconstriction and/or lung composition, altergies and/or bronchoconstriction and/or lung complianmation, altergies, asthma, impeded respiration, respiratory composition such as pulmonary vasoconstriction are associated conformation, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonacleotides into products that free adenosine into the system content of the serves to prevent the breakdown of the oligonacleotides into products that free adenosine into the system content of the serves to prevent the breakdown of the oligonacleotides into products that free adenosine into the system content of the online and thereby, to prevent any unwanted effects due to it
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel composition (a) a first active comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies reducing adenosine sensitivity, levels of adenosine (A) or (A)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
 AA281534-derived oligonucleotide SEQ ID 3687.
                                     29-JUL-2004
                                                                        ABD24675;
                                                                                                         ABD24675
                                                                                                                                                                                                                                                                                                                                              Sequence
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L, Shahabuddin
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Query Match Best Local : Matches 1:

Similarity

88.9%;

Score 14.8; Pred. No. 4.

3; DB 1; 4.4e+02;

Length 20;

Indels

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Gaps

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Conservative

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CC oligonuclectides are derived from a gene encoding or regulating manual dysfunction or cancer and can be anti-sense to the corresponding mana. CC device, in separate containers, (b) that comprises: (a) a delivery construction or cancer and can be anti-sense to the corresponding mana. CC device, in separate containers, (b) the oligonucleotides, (c) anstructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analysis of treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mana or conformation, and/or bronchoconstriction and/or lung configuration, and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, respiratory configuration, allergies, asthma, impeded respiration, respiratory transplantation rejection, pulmonary infections, bronchitis or cancer. C the induced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system content any unwanted effects due to it
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Miller
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C, Shahabuddin
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CC comprising oligonucleotides, effective for alleviating composition of ligonucleotides, effective for alleviating compositions adenosine sensitivity, levels of adenosine (A) or (A) receptors, composition or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating context or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery construction for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenargic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition composition composition of reduce the amount of target polypeptide present in the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition and/or burndary obstruction, and/or bronchoconstriction and/or lung cinflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as perjaration, respiratory disease, pulmonary condition, allergies, asthma, impeded respiration, respiratory condition, emphysema, chronic obstructive pulmonary disease, pulmonary disease, pulmonary conditions but a constitution and or analytic reference and conditions of the condition are associated constructive pulmonary disease, pulmonary disease, pulmonary conditions but a condition such as a condition and condition are associated conditions and conditions of the condition are associated conditions are associated conditions and conditions and conditions are associated conditions. The condition are conditions are associated conditions are associated conditions are associated conditions.
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   transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding the the sense oligos corresponding to the manual present in the target RNA serves to prevent the breakdown of the care of the content of the care of the c
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comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonuclectides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered
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Pred. No. 4.4e+02
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RESULT 76
ABD22850
ID ABD22850
AC ABD2
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XX ABD2
XX Huma
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Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction;
This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and
                                                                                                                                                                                                             Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                         Claim 15; SEQ
                                                                                                                                                                                      bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                              Miller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human myosin X-derived oligonucleotide SEQ
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16; Conserv
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, Tang L,
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                                                                                                                         ID NO 1862; 763pp; English
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L, Shahabuddin
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Pred. No. 4.4e+02
0; Mismatches
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ABD24496
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RESULT 761
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Best Local S
Matches 16
                Nyce JW,
Miller S
                                                                                                                                                                                                                                                                                                                                                 Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic,
                                                                                                                                                                                                                                                                                                              emphysema, chronic obstructive pulmonary disease; cancer; bronchitis,
pulmonary transplantation rejection; ss; primer.
                                                                                                                24-APR-2001; 2001US-0286036P
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                                                                           (EPIG-)
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              Li Y,
Tang
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                                                                                                                                                       2002WO-US013143
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                Sandrasagra A,
L, Shahabuddin
                                                                           PHARM INC
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88.9%;
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Pred. No. 4.
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                                    Katz E,
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                                    Pabalan J,
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RESULT 762
ABD25532
ID ABD255
XX ABD255
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XX 11256
DT 29-JUL
DT 29-JUL
CX ALI256
XX Human,
KW Fespir
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                                                                                                                                                             Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           AI125651-derived oligonucleotide SEQ ID 4544.
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   WO200285309-A2
                                                                     Homo sapiens.
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                                                                                                                                           pulmonary transplantation
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16; Conser
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                                                                                         CC comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and CC reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, CC surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating CC expression of a target polypeptide associated with lung airway or lung CC dysfunction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered CC composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The CC pulmonary obstruction, and/or bronchoconstriction and/or lung CC inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, comprises oligonary obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary hypertension, emphysema, chronic obstructive pulmon, bronchitis or cancer.
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The reduced adenosine content of the anti-sense oligos
thymidines present in the target RNA serves to prevent
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CC This invention describes a novel composition (a) a first active agent, CC comprising oligonucleotides, effective for alleviating CC bronchoconstriction, respiratory tract inflammation, allergies and CC reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, CC surfactant depletion or hyposecretion, when administered to a mammal. The CC oligonucleotides are derived from a gene encoding or regulating CC expression of a target polypeptide associated with lung airway or lung CC dysfunction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) containers, (b) the oligonucleotides, (c) containers, (b) the oligonucleotides, (c) containers, (c) the invention has antiallergic, antiinflammatory, antiasthmatic, CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a CC treating a respiratory, lung or malignant disease. The administered CC composition comprises oligo and is administered to reduce the production
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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptor
                                                                                                                                                                                                                           Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
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RESULT 767
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC dystunction or cancer and can be anti-sense to the corresponding mayar. CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, and the invention has antiallergic, antiinflammatory, antiasthmatic, and the addrenergic agonist. The composition is useful for preventing or composition composition is useful for preventing or composition comprises oligo and is administered to reduce the production of treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mana or to composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mana or to composition, allergies and/or bronchoconstriction and/or lung complete the amount of target polypeptide present in the lungs. The composition action, allergies and/or surfactant hypoproduction are associated composition, allergies, asthma, impeded respiration, respiratory composition, allergies, asthma, impeded respiration, respiratory composition, emphysema, chronic obstructive pulmonary disease, pulmonary composition, emphysema, chronic obstructive pulmonary disease, pulmonary composition, emphysema, chronic obstructive pulmonary disease, pulmonary composition, pulmonary of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system content and emphysema chronic disease composition in the target mana composition into the system composition and effects due to it.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local &
                                                  Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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     WPI; 2003-093058/08
                                                                                                                                                                                   24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                      23-APR-2002; 2002WO-US013143
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                                                                                                                                   EPIGENESIS PHARM INC
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                                                  Li Y,
Tang
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                                                     Sandrasagra A,
L, Shahabuddin
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Pred. No. 4.4e
0; Mismatches
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                                                                                Katz
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                                                                                Pabalan
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This invention describes a novel composition (a) a first active agent, comprising oligonuclectides, effective for alleviating troubconstriction, respiratory tract inflammation, allergies and composition adenosine sensitivity, levels of adenosine (A) or (A) receptors, containing adenosine sensitivity, levels of adenosine (A) or (A) receptors, contained to the corresponding or requiating controllection or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating mRNA. Contained to a describe a kit, that comprises: (a) a delivery contained to the invention or cancer and can be anti-sense to the corresponding mRNA. Contained to the invention has antiallergic, antiinflammatory, antiasthmatic, confit the invention has antiallergic, antiinflammatory, antiasthmatic, canadgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition of composition comprises oligo and is administered to reduce the production corrected the amount of target polypeptide present in the lungs. The comparison of the amount of target polypeptide present in the lungs. The comparison and/or bronchoconstriction and/or lung confilmmation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, and/or bronchoconstriction and/or lung inflammation, allergies, asthma, impeded respiration, respiratory condition rejection, pulmonary infections, bronchitis or cancer. Continually present in the target mRNA or to the oligonucleotides into products that free adenosine into the breakdown of the oligonucleotides into products that free adenosine into the system condition and prevent any unwanted effects due to it
  Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targe nucleic acids associated with lung airway or lung dysfunction,
Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 15;
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  BP;
20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                               Matches
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Best Local (
              1520 АААААААААААСТАААА 1537
                               16;
μ.
                                       Similarity
AAAAAAAAAAAAAAAAA
                                Conservative
                                      1.1%;
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                                               Score 14.8;
                                       Pred. No.
                                Mismatches
                                       4.4e+02
                                                DB 1;
                               2
                                               Length 20
                               0
                               Gaps
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ABD25111 standard; DNA; 20
                              ABD25111;
                                 ВP
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AI125228-derived 29-JUL-2004

oligonucleotide SEQ ID

4123.

(first entry)

Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.

Homo sapiens.

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RESULT 769
ADH08684
ID ADH086
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC comperising oligonucleotides, effective for alleviating CC reducing adenosine sensitivity, levels of adenosine (A) receptors, CC surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating CC expression of a target polypeptide associated with lung airway or lung CC dysfunction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition CC instructions for adding a carrier and for use of the kit. The composition containers, (b) the oligonucleotides, (c) analysesic, hypotensive, immunosuppressive and cytostatic activity, is a beta advancer; agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered CC composition comprises oligo and is administered to reduce the production or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The CC inflammation, allergies and/or bronchoconstriction and/or lung ulmonary obstruction, and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, emphysema, chronic obstructive pulmonary disease, pulmonary CC thymidines present in the target RNA perves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system of the oligonucleotides into products that free adenosine into the system of the oligonucleotides into products that free adenosine into the system of the content of the surfaction and thereby, to respect the oligonucleotides into products that free adenosine into the system of the content of the anti-sense oligonucleotides into force the content of the surfaction of the content of th
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Best Local
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Linking oligonucleotide; 88; nucleic acid detection;
                                             Nanotechnology nucleic acid detection method associated
                                                                                                  11-MAR-2004
                                                                                                                                                                                             ADH08684 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               prevent any unwanted effects due to it
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, Tang L,
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L, Shahabuddin
                                                                                                                                                                                             DNA;
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Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                            8;
4.4e+02;
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Matches
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Best Local :
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25-JUN-1999;
26-APR-2000;
26-JUN-2000;
                                                                                                                                                                                Linking oligonucleotide; ss; nucleic acid detection; nanoparticle-oligonucleotide conjugate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detecting nucleic acid used for e.g. diagnosis of diseases, fo DNA sequencing, comprises observing detectable change caused by hybridization of nucleic acid with substrate or particle bound
12-OCT-2001; 2001US-00976617.
                                             26-SEP-2002.
                                                                                           US2002137072-A1
                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                   Nanotechnology nucleic
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; 99US-00344667.
; 2000US-0200161P.
; 2000US-00603830.
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                                                                                                                                                                                                                                                                                                    (first entry)
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97WO-US012783
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                                                                                                                                                                                                                                                        acid detection method associated
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RESULT 771
ADH0874
ID ADH087
XX ADH087
XX ADH087
XX I1-MAR
XX Linkin
KW Linkin
KW Inanopa
XX Synthe
XX US2002
PD 26-SEE
XX 10-OCT
XX 29-JUI
PR 29-JUI
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Best Local S
Matches 16
                                          29-JUL-1996;
21-JUL-1997;
29-JAN-1999;
25-JUN-1999;
26-APR-2000;
26-JUN-2000;
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21-JUL-1997;
29-JAN-1999;
25-JUN-1999;
26-APR-2000;
26-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a method of detecting a nucleic acid with at least two portions by providing a type of nanoparticle-oligonucleotide conjugate, contacting the nucleic acid and nanoparticles to allow hybridisation of the oligonucleotides with the two or more portions of the nucleic acid and observing a detectable change brought about by hybridisation. The oligonucleotides have a sequence complementary to the sequence of at least two portions of the nucleic acid. Hybridisation of the oligonucleotides on the nanoparticles with the nucleic acid results in a detectable change. This sequence represents a linking oligonucleotide of the invention.
                                                                                                                                                                                                              10-OCT-2001; 2001US-00974007.
                                                                                                                                                                                                                                                           26-SEP-2002
                                                                                                                                                                                                                                                                                                      US2002137071-A1
                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                           Linking oligonucleotide; ss; nanoparticle-oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nanotechnology nucleic acid detection method associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-MAR-2004
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    (NANO-)
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16; Conserv
       NANOSPHERE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP;
                                             99US-00240755.
99US-00344667.
2000US-0200161P.
2000US-00603830.
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2000US-00603830.
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97WO-US012783.
99US-00240755.
99US-00344667.
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                                                                                                                                                                                                                                                                                                                                                                                              nucleic acid detection; conjugate.
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Pred. No. 4.
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Mismatches
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4.4e+02;
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RESULT 772
ADH65941/c
ID ADH659
XX ADH659
XX ADH659
XX ADH659
XX ADH659
XX ATISE
KW ANTISE
KW INFIAM
CARTIC
KW PHOMO E
XX HOMO E
YX WO2003
XX HOMO E
YX WO2003
XX CO-MA1
XX PT CO-MA1
XX PT CARTÍC
XX PF CO-MA1
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Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetes; obseity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detecting nucleic acid used for e.g. diagnosis of diseases, forensics and DNA sequencing, comprises observing detectable change caused by hybridization of nucleic acid with substrate or particle bound
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                                                                                                                                                                                                                                                                         Crosby SD,
                                                                                                                                                                                                                                                                                                                                                                                     20-MAY-2002;
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                                                                                                                                                                                                                                                                                                                               (PHAA ) PHARMACIA CORP
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Pred. No. 4
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New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, ob cardiovascular disorder, hyperlipidemia or Cushing's syndrome.

Claim

4; SEQ

IJ

985pp; English

comprises NO 2775;

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antisense oligonucleotides

that

are

targeted

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RESULT 773
ADH67409/c
ID ADH674
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Matches
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                                                                                                                                                                                                                                    The invention comprises an antisense oligonucleotides that are targeted to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obseity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense migonucleotide that targets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.
                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; SEQ ID NO 4243; 985pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-035034/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-MAY-2002; 2002US-0381857P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-MAY-2003; 2003WO-US016084.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003099215-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense oligonucleotide; glucocorticoid receptor; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human glucocorticoid receptor-specific antisense oligonucleotide #4243.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADH67409
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PHAA ) PHARMACIA CORP
Local Similarity
mes 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                               human glucocorticoid
tains 2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1522 AAAAAAAAAAAGTAAAAGG 1539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   g,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
                                                                                                                      20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAAGAAAAAAATAAAAGG 2
                                                                                                                      ₽P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nalseth AE;
      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                   1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 C; 1 G; 15 T; 0 U; 0 Other;
                                                                                                                      1 C; 1 G; 17 T; 0 U; 0 Other;
                               88.9%;
                                                        1.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                               receptor gene. NOTE: The present sequence (2'-MOE) wings and a phosphorothioate bac
<u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                            Score 14.8; DB 1; Length 20; Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8; DB 1;
Pred. No. 4.4e+02;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20;
0,
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                                                                                                                                                                                  backbone
Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
0
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RESULT 775 ADI47212

ADI47212 standard; DNA; 20

ВP

HXXXX

ADI47212;

22-APR-2004

(first entry)

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RESULT 774
ADI34492
ID ADI344
XX ADI344
XX Nucleo
XX Nuclei
XX Nuclei
XX Synthe
XX Synthe
XX Synthe
XX Synthe
XX HO2003
XX II-DEC
XX SYNTH
XX II-DEC
XX SYNTH
XX II-DEC
XX SYNTH
XX SYNTH
XX SYNTH
XX SYNTH
XX SYNTH
XX SYNTH
XX II-DEC
XX SYNTH
XX II-DEC
XX II-DEC
XX III-DEC
XX 
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                                                                                                                                                                                                                                                                                                                                            The invention relates to amplifying for RNA in a sample comprises caliminating single-stranded oligonuclectide from the transcription sample. The method involves synthesizing single-stranded cDNA by incubating the sample RNA with reverse transcriptase and an of the RNA; converting the single-stranded cDNA into double-stranded cDNA to form a transcription sample containing a cDNA template; eliminating single-stranded cDNA template; eliminating the form the transcription sample; and transcribing the cDNA template into RNA using an RNA polymerase. The method is useful for improving RNA polymerase based RNA transcription from a polymucleotide template. The method inhibits the undesired non-template derived production of RNA in the transcription reaction. The present sequence represents an oligonucleotide used to exemplify RNA transcription in the presence of single- and double-stranded
                                                                                                                               Matches
                                                                                                                                                          Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Amplifying for RNA in a sample, useful for improving RNA polymerase based RNA transcription from a polynucleotide template, comprises eliminating single-stranded oligonucleotide from the transcription sample.
                                                                                                                                                                                                                                                            Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-MAY-2002; 2002US-0384454P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid
                                                                                                                                                                                                                                                                                                                        oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; SEQ ID NO 11; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kamme FC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-MAY-2003; 2003WO-US017103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2003102243-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADI34492;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADI34492 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (JANC ) JANSSEN PHARM NV
                                                               1520 AAAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1521 ААААААААААСТААААС 1538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                               16;
1 AAAAAAAAAAAAAAAAA 18
                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAAAAAAAAAAAAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zhu JY;
                                                                                                                                                                                                                                                        BP;
                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amplification; RNA transcription; RNA polymerase; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                        20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                       1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ₽₽
                                                                                                                            <u>.</u>
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                                                                                                                               Pred. No. 4.4e+02
0; Mismatches
                                                                                                                                                                                        Score 14.8;
                                                                                                                                                                                               DB 1;
                                                                                                                               Indels
                                                                                                                               ٥,
                                                                                                                            Gaps
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RESULT 776
ADJ51142/c
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                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a novel method involving forming a laminar flow, by passing into a micro channel, a solution containing the specimen molecules, and a solution containing probe molecules capable of forming a complex with the specimen molecules. The dispersion of the formed complex is selectively promoted, based on their affinity, and the degree of dispersion of the complex formed between the specimen molecules and the probe molecules is detected and analysed. The probe molecules are capable of producing fluorescence. This polymucleotide sequence represents an oligo used in the exemplification of the invention.
                                                                            modified_base
                                                                                                                    Synthetic
                                                                                                                                                                                    Polyalkyleneamine-conjugated oligonucleotide #1.
                                                                                                                                                                                                                 06-MAY-2004
                                                                                                                                                                                                                                            ADJ51142;
                                                                                                                                                                                                                                                                   ADJ51142
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 9; 19pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  forming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Analysis of sample molecules such as DNA fragment, by using micro channel to form laminar flow of specimen molecule-containing solution and complex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-180318/17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yamashita K,
Yamaguchi Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-JUL-2002; 2002JP-00211462.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-JUL-2003; 2003WO-JP009142.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2004010140-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                laminar flow; micro channel; complex; selectively promoted; fluorescence;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Molecule analysing
                                                                                                                                              inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NAAD-)
                                                                                                                                                            Antimicrobial; Antiinflammatory; Cytostatic; prodrug; infection;
                                                                                                                                                                                                                                                                                                                                                                  1520
                                                                                                                                                                                                                                                                                                                                       \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  molecule containing solution.
                                                                                                                                                                                                                                                                      standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                               ААААААААААСТААА 1537
                                                                                                                                                                                                                                                                                                                                       ААЛАААААААААААА 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INST
                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Maeda H,
                                                                                                                                                tumour.
           /mod_base= OTHER
/note= "Optionally conjugated with spermine,
polyethylenimine (PEI) 600 or PEI 1200,
tetraethylenpentamine. Also optionally 5'-pr
                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADVANCED IND SCI & TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             microchannel method related probe
                                                                                                                                                                                                                                                                                                                                                                                                      1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shimizu H,
                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8; DB 1;
Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Miyazaki M,
                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                   Length
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                                                                                                                                                                                                                                                                                                                                                                                            Indels
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             5'-protected
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               with
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RESULT 777
ADK99410/c
ID ADK994
XX
AC ADK984
XX
DT 06-MAY
XX
DE Primer
XX
KW human;
XX
COS Synthe
XX
Synthe
XX
AX
DF2003
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     fusogenic part is a lipophilic polyamine, polyethylenimine, polyalylamine, fusogenic peptide, oligomeric imidazole, histidine, pyridine, hydroxylamine, substituted hydroxylamine, hydrazine, pyridine, hydrazine, thiourea or imine. The targeting part is a ligand that binds to a cellular reporter, where the targeting part is that binds to a cellular reporter, where the targeting part is transferrin, folate, epidermal growth factor, nerve growth factor, insulin, alpha-fetoprotein, galactose, galactosamine, lactose, mannose, a polyclonal antibody, monoclonal antibody, vitamin B12, bbuprofen, cholesterol, low-density lipoprotein, peptide comprising an arginine-conclusional antibody and oligonucleotide analogue, a peptide nucleic acid or a peptide nucleic acid sequence. The oligomeric part is an oligonucleotide, and oligonucleotide analogue, a peptide nucleic acid or a peptide mucleic acid analogue. OC is useful as a prodrug, useful in capaniams. The present sequence represents an oligonucleotide used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                part, a fusogenic part, and a targeting part; and enhancing the cellular uptake of OC, by conjugating OC to a fusogenic part. In (C), the fusogenic part is covalently linked to the oligomeric part. The targeting part is covalently linked to the oligomeric or fusogenic part, where the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel polyalkyleneamine-containing oligomeric compound useful for preventing or delaying infection, inflammation or tumor formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a polyalkyleneamine-containing oligomeric compound (OC). Also described is a compound (C) comprising an oligon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Manoharan M, Guzaev AP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-JUL-2002; 2002US-00199585
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                                                                                         human; single nucleotide polymorphism;
                                                                                                                              Primer of the
                                                                                                                                                                 06-MAY-2004
                                                                                                                                                                                                    ADK98410;
                                                                                                                                                                                                                                        ADK98410 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3; Page 22; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MANO/) MANOHARAN M. (GUZA/) GUZAEV A P. (MAIE/) MAIER M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-JUL-2002; 2002US-00199585
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                                                                                                                                                                                                                                                                                                                                                                1520 ААЛАЛАЛАЛАЛАСТАЛАЛ 1537
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                                                                                                                                                                                                                                                                                                                             20 AAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                     16;
                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the invention.
                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                 (first entry)
                                                                                                                              invention #4130.
                                                                                                                                                                                                                                                                                                                                                                                                                      88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. 4.4e+02
                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                            SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                            88;
                                                                                                                                                                                                                                                                                                                                                                                                     2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          comprising an oligomeric
                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 20
                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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JP2003259875-A. Synthetic

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ARESULT 778
ADJ60935/c
ID ADJ609
XX ADJ609
XX O6-MAY
XX O1igon
XX Interl
XW interl
XW cystic
XX Homo s
XX 
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a polynucleotide isolated from a gene and is useful for detecting a single nucleotide polymorphism human gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The present sequence represents a primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome;
                                                                       Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
                                                                                                                                                                                                         WPI; 2004-203534/19.
                                                                                                                                                                                                                                                           Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                   29-JUL-2002; 2002US-0399076P
                                                                                                                                                                                                                                                                                                                                                                                                                                        25-JUL-2003; 2003WO-US023509
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2004011613-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide associated to PDE4C #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADJ60935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel polynucleotide useful for PCR amplification along with two DNA fragment from another set of sequences, or for detecting single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-093977/10.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleotide polymorphism in human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-SEP-2003
                                                                                                                                                                                                                                                                                                                                      (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-MAR-2002; 2002JP-00064373
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
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16; Conserv
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                                                                                                                                                                                                                                                      Tang L,
lin S, Lu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CTCACCTCTGAGGCCTCT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 7439; 2627pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 5 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                      Sandrasagra
H, Cong H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.8; D
Pred. No. 4.4e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                ۶
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene.
                                                                                                                                                                                                                                                                              Aguilar D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.40+02;
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                                                                                                                                                                                                                                                                                      Miller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               from a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ä
                                                                                                                            e.g.,
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SEQ

ID NO 1791; 85pp; English

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RESULT 779
AD132920
ID AD1329
XX AD1329
XX AD1329
XX O1190
XX INAMPLE AD1329
XX INAMPL AD13
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Best Local (
                                                                                                                    28-MAR-2001;
09-APR-2001;
10-AUG-2001;
09-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                              28-MAR-2000;
26-APR-2000;
26-JUN-2000;
26-JUN-2000;
                                                                                                                                                                                                                   11-DEC-
11-DEC-
12-JAN-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease is chosen from airway inflammation, allergy(les), asthma, impeded respiration, cystic fibrosis (CP), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nanoparticle; cell line auth
                           (PARK/)
(TATO/)
                                                                                                                                                                                                                                                                                            08-DEC-2000;
                                                                                                                                                                                                                                                                                                                   08-DEC-2000;
                                                                                                                                                                                                                                                                                                                                          11-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-JUN-1999;
13-JAN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-OCT-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-NOV-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligo related to thiol oligo-gold colloid conjugate probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
     (MIRK/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-JUL-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1520 ААЛАЛАЛАЛАЛАГТАЛАЛ 1537
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                                                                                                                                                                                                                   -2001;
                                                                                                                                                                                                                                                                 -2000;
) PARK S.
) TATON T A.
) MIRKIN C A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        :le; gold; disease; forensic; paternity authentication; gene therapy; ss; gold
                                                                                             2000US-0176409P

2000US-012699P

2000US-0200161P

2000US-02001390P

2000US-0224631P

2000US-0254392P

2000US-0254392P

2000US-025431SP

2000US-0255236P

2000US-0255236P

2001US-00760500

2001US-00820279

2001US-0282640P

2001US-0282640P

2001US-03278640P

2001US-03278640P

2001US-03278640P

2001US-03278640P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2002US-00266983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-00240755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97WO-US012783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96US-0031809P
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G;
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Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           testing;
colloid conjugate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ
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RESULT 780
AD132905
ID AD1329
XX AD1329
XX AD1329
XX O6-MAY
XX Synthe
XX nanopa
KW cell l
KW probe.
XX O6-NOV
XX O6-NOV
XX O8-OC1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       having at least two portions comprising contacting the nucleic acid with at least two types of nanoparticles, such as gold, having attached coligonucleotides and observing a detectable change brought about by hybridisation of the oligonucleotides on the nanoparticles with the nucleic acid. The method of the invention may be useful for detecting a nucleic acid, preferably viral RNA or DNA, bacterial DNA, a gene associated with a disease, a fungal DNA, synthetic DNA or RNA, a gene cutilised chain reaction amplification. The detected nucleic acid may be utilised for diagnosis of disease, sequencing of nucleic acids, forensics, paternity testing, cell line authentication and monitoring gene therapy. The method for detecting the nucleic acids is based on observing a colour change with the naked eye and is cheap, fast, simple, and requence is that of the oligonucleotide which is related to a thiol-modified oligonucleotide-gold colloid conjugate probe of the invention.
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Best Local S
   29-JUL-1996; 96US-0031809P.
21-JUL-1997; 97WO-US012783.
29-JAN-1999; 99US-00240755.
25-JUN-1999; 99US-00344667.
13-JAN-2000; 2000US-0176409P.
28-MAR-2000; 2000US-0122699P.
26-APR-2000; 2000US-0200161P.
26-JUN-2000; 2000US-02013906P.
26-JUN-2000; 2000US-0213906P.
11-AUG-2000; 2000US-0224631P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Detection of nucleic acid, e.g. viral RNA or DNA, comprises contacting nucleic acid with different types of nanoparticles having attached oligonuclectides and observing detectable change brought about by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Park
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hybridization.
                                                                                                                                                                                                                                                                                                                                                         08-OCT-2002; 2002US-00266983
                                                                                                                                                                                                                                                                                                                                                                                                                       06-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US2003207296-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nanoparticle; gold; disease; forensic; paternity testing;
cell line authentication; gene therapy; ss; gold colloid conjugate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic thiol-modified oligo-gold colloid conjugate probe -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADI32905;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADI32905 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 24; SEQ ID NO 70; 206pp; English.
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16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAAAAAAAAAAAAAAA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Taton TA, Mirkin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.1%;
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Pred. No. 4.4e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.4e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
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RESULT 781 ADK70840

ADK70840 standard;

DNA;

20 ВP 맑 á

1520 AAAAAAAAAAAGTAAAA 1537

1 AAAAAAAAAAAAAAAA 18

Query Match Best Local (Matches

16;

Conservative

0;

Similarity

1.1%;

Score 14.8; Pred. No. 4. Mismatches

4.4e+02;

2

<u>,,</u>

Gaps

0

DB 1; Length 20

06-MAY-2004

(first entry)

ADK70840;

DNA preparation; 5' mRNA; linker synthesis; gene regulation; gene expression; ss; tag.

primer synthesis;

WO2003106672-A2 Unidentified 5' mRNA DNA preparation method related tag DNA sequence #8

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                                                                                                                 CC having at least two portions comprising contacting the nucleic acid with CC at least two types of nanoparticles, such as gold, having attached CC oligonuclectides and observing a detectable change brought about by CC hybridisation of the oligonuclectides on the nanoparticles with the CC nucleic acid. The method of the invention may be useful for detecting a CC nucleic acid, preferably viral RNA or DNA, bacterial DNA, a gene CC associated with a disease, a fungal DNA, barthetic DNA or RNA, CC structurally modified natural or synthetic DNA or RNA or a product of a polymerase chain reaction amplification. The detected nucleic acid may be CC utilised for diagnosis of disease, sequencing of nucleic acids, CC utilised for diagnosis of disease, sequencing of nucleic acids, CC gene therapy. The method for detecting the nucleic acids is based on CC observing a colour change with the naked eye and is cheap, fast, simple, CC and robust, requiring no specialised or expensive equipment. The current sequence is that of the synthetic thiol-modified oligonucleotide-gold CC colloid conjugate probe of the invention which is linked via a thiol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-DEC-2000; 2000US-0254392P.
08-DEC-2000; 2000US-0254318P.
11-DEC-2000; 2000US-0255235P.
11-DEC-2000; 2000US-0255236P.
12-JAN-2001; 2001US-00760500.
26-MAR-2001; 2001US-00820279.
09-APR-2001; 2001US-0282640P.
10-AUG-2001; 2001US-00327777.
09-OCT-2001; 2001US-0008978.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detection of nucleic acid, e.g. viral RNA or DNA, comprises contacting nucleic acid with different types of nanoparticles having attached oligonucleotides and observing detectable change brought about by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (PARK/)
(TATO/)
Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                      group to a gold nanoparticle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a novel method for detecting a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 18; SEQ ID NO 55; 206pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hybridization.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-059754/06.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MIRK/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MIRKIN C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PARK S.
TATON T A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Taton TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mirkin
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RESULT 782
ADK69880/c
ID ADK698
XX ADK698
XX O6-MAY
XX Phosph
XX P
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      corresponding to a nucleotide sequence of a 5' end of an mRNA. The method is useful for synthesising a nucleotide sequence to be used as a linker or primer and selectively collecting multiple nucleic acid fragments containing information on the nucleotide sequences at the 5' end of multiple mRNA in a sample. The method is also useful for identifying regions in the genome, which are required for gene regulation and gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Preparing DNA fragment corresponding region of mRNA, by preparing nucleic sequence of 5' end of mRNA, cleaving
                                                     11-JAN-2000; 2000US-00481486.
10-JAN-2001; 2001WO-US000715.
                                                                                                                                                                                   13-NOV-2003.
                                                                                                                                                                                                                                      US2003212267-A1.
                                                                                                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Phosphorothioate backbone; sulphurised oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sulphurised oligonucleotide #10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADK69880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADK69880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 0 A; 1 C; 10 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   expression.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIKE )
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12-AUG-2002; 2002JP-00235294.
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                                                                                                                                12-DEC-2002; 2002US-00181200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-JUN-2003; 2003WO-JP007514.
     (COLE/) COLE D L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               545 rerecerecerecere 562
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16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               rerererererecere 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present DNA sequence was used in an example of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           즛
                                                                                                                                                                                                                                                                                                                                                            /*tag=
                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
/note= "Phosphorothioate backbone; 2'-0-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                         residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Harbers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               to nucleotide sequence of 5' end acid corresponding to nucleotide nucleic acid with restriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
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RESULT 783
ADX69885/c
ID ADX698
XX ADX698
XX Sulphu
XX Phosph
XX Phosph
XX Uniden
XX Uniden
XX Uniden
XX US2003
XX I3-NOV
PF 12-DEC
XX I1-JAN
PR 10-JAN
XX (COLE,
PA (RAVI,
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  phosphite intermediate; and oxidising the phosphite intermediate with an acetyl disulfide in an acetonitrile for a time to effect conversion of the phosphite intermediate to phosphorothioate. The invented method achieves high yields and greater efficiency. The present sequence is sulphurised oligonucleotide used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Preparation of phosphorothioate oligonucleotides involves oxidizing phosphite intermediate with acetyl disulfide in acetonitrile for ti effect conversion of phosphite intermediate to phosphorothioate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to phosphorothioate oligonucleotides having nucleoside with 240 modification are prepared by phosphitylating 5'-hydroxyl of a nucleic acid moiety having a nucleoside with 2' modification in an acetonitrile containing solvent mixture to form a
  Preparation of phosphorothioate oligonucleotides involves oxidizing
                                                     WPI; 2004-069376/07.
                                                                                                                                                                                                                                                             11-JAN-2000;
10-JAN-2001;
                                                                                                                                                                                                                                                                                                                                        12-DEC-2002; 2002US-00181200
                                                                                                                                                                                                                                                                                                                                                                                                                                                 US2003212267-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Phosphorothicate backbone; sulphurised oligonucleotide; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sulphurised oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 12; SEQ ID NO 10; 8pp; English.
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                                                                                                         Cole DL,
                                                                                                                                                      (COLE/) COLE D L.
(RAVI/) RAVIKUMAR V T.
(CHER/) CHERUVALLATH Z
                                                                                                                                                                                                                                                                                                                                                                                           13-NOV-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unidentified
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2001WO-US000715.
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/note= "Phosphorothioate backbone; 2'-O-methoxyethyl
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0; Mismatches
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Pred. No. 4.4e+02;
                                                                                                         SZ
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RESULT 784
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XX Chimer
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                          associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate pain from burns, migraine headache, cluster headache, mild-to-moderate pain from burns, migraine headache, cluster headache, mild-to-moderate pain fastilahood seizure disorder, including headache, seizure disorder such as childhood seizure disorder, including
                                                                                                                                                                                         The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition
                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                       Claim 4; SEQ ID NO 1981; 417pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-203785/19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Navl.3; Analgesic; Nootropic; Neuroprotective; diabetic neuropathy; arthritic pain; migraine l
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Pred. No. 4.4e+02;
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headache;
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RESULT 785
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Best Local S
Matches 16
                                                                                   nucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, lower back pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with 2 MOE wings and a deoxy gap. Used during the antisense inhibition of human Nav1.3 expression, the oligonucleotides are designed to target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence represents a chimeric phosphorothicat 2'MOE wings and a deoxy gap. Used during the a human Navl.3 expression, the oligonucleotides different regions of the human Navl.3 RNA.
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Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to an antisense compound targeted nucleic acid molecule encoding Nav1.3, where the antisense compo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nav1.3; Analgesic; Nootropic; Neuroprotective; diabetic neuropathy; arthritic pain; migraine l
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      BP;
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                                                                of the human Nav1.3 RNA
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   1 C; 0 G; 19
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Pred. No. 4
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Query Match Best Local S Matches 16

Similarity

1.1%;

Score 14.8; DB 1 Pred. No. 4.4e+02 0; Mismatches

DB 1;

Length 20;

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Gaps

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RESULT 786
ADK74969/c
ID ADK749
XX ADK749
XX ACAMAY
XX Chimer
XX Navl.3
XW diabet
XX Synthe
XX Synthe
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RESULT 787
ADK75847
ID ADK758
XX
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Best Local S
Matches 16
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                             ADK75847 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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dlarity 88.9%;
Conservative
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RESULT 788
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Best Local
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                           Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache;
                                                                                                                                                                           ADK79524 standard; DNA; 20
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                                                                               Chimeric
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                                                                            phosphorothioate oligonucleotide to target Nav1.3 #6858.
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               epilepsy; ataxia;
                                                                                                                                                                                                                                                                                                                             Conservative
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Pred. No. 4.4e+02;
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Matches 16
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                              14-AUG-2002; 2002US-0403416P
                                                                         14-AUG-2003; 2003WO-US025465
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                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                  Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic diabetic neuropathy; arthritic pain; migraine headache;
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                                                                                                                                                                                                                                                                           infantile epilepsy; ataxia; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 BP; 11 A; 1 C; 2 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TTTTGATTTTTAATCACA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             relates to an antisense compound targeted encoding Nav1.3, where the antisense compo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8; DB 1;
Pred. No. 4.4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     condition associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                            neuralgia;
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RESULT 790
ADK74889/c
ID ADK748
XX ADK748
XX ADK748
XX Chimex
XX Nav1.3
KW Giabet
KW infant
XX Synthe
XX WO2004
XX WO2004
XX II -AUC
XX II -AUC
XX PHAA
XX PFI Roberc
XX WPI;
XX PFI Nav1;
PFI With!
PFI With!
PFI Wisor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, chapter in the pain from burns, migraine headache, cluster headache, mild-to-moderate pain from burns, migraine headache, cluster headache, mild-to-moderate pain neuropathy surfaine headache, cluster headache, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with 2°MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target odifferent regions of the human Navl.3 RNA.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                          Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; SEQ ID NO 2022; 417pp; English.
                                                                                                                                                                                                          14-AUG-2002; 2002US-0403416P
                                                                                                                                                                                                                                             14-AUG-2003; 2003WO-US025465.
                                                                                                                                                                                                                                                                                                                    WO2004016754-A2
                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chimeric phosphorothioate oligonucleotide to target Nav1.3 #2223.
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                                                                                                                                                                    (PHAA)
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                                                                                                                                                                      PHARMACIA CORP
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Pred. No. 4.4e+02;
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Claim

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SEQ ID

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2223; 417pp; English

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RESULT 791
ADK80788/c
ID ADK807
XX ADK807
XX ADK807
XX ADK807
XX ADK807
XX Chimer
XX Navl.3
KW diabet
KW infant
XX WO2004
XX WO2004
XX IA-AUG
XX (PHAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cc nucleic acid molecule encoding Navl.3, where the antisense compound CC specifically hybridizes with and inhibits the expression of Navl.3. The CC compound and composition are useful for treating a disease or condition CC associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, CC diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, CC pain from burns, migraine headache, cluster headache, mild-to-moderate CC headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present Sequence represents a chimeric phosphorothicate oligonucleotide with CC 2'MOS wings and a deoxy gap. Used during the antisense inhibition of CC human Navl.3 expression, the oligonucleotides are designed to target XX different regions of the human Navl.3 RNA.
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Best Local S
Matches 16
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   The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound specifically pybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pair diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain,
                                                                                                                                                                                                                                                                                                New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-AUG-2002;
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                                                                                                                                                                                                                                            SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
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nilarity 88.9%;
Conservative
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                                                                                                                                                                                                                                     NO 8122; 417pp; English
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Pred. No. 4.4e
0; Mismatches
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                                 lower back pain,
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                           The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound generifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, cliabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate pain from burns, migraine headache, cluster headache, mild-to-moderate pain neuropathy, trigeminal neuropathy, arthritic pain, acute pain, cheadache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target different regions of the human Navl.3 RNA.
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Best L
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                                                                                                                                                                                                                                                                                                                            New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                              Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diabetic neuropathy; arthritic pain; migraine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chimeric phosphorothioate oligonucleotide to target Navl.3 #2172
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nav1.3; Analgesic; Nootropic; Neuroprotective;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (PHAA ) PHARMACIA CORP
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                                                                                                                                                                                                                                                                                              SEQ ID NO 2172; 417pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8;
Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cluster headache, mild-to-moderate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       멂
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     headache
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   post-herpetic neuralgia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                         lower back pain,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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ADX7721
ADX772
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 793
                                                                                                                                                                                                                                                                                     The present invention relates to an antisense compound targeted to a concleic acid molecule encoding Navl.3, where the antisense compound compound hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, compound in from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target of different regions of the human Navl.3 RNA.
                                                                                                                                               Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-AUG-2002; 2002US-0403416P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chimeric phosphorothioate oligonucleotide to target Nav1.3 #4552
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (PHAA ) PHARMACIA CORP.
                                                           1165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2004-203785/19
                                                                                                                      16;
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16; Conserv
   N
                                                                                                                                               Similarity
                                                                                                                                                                                                                                      20
                                                           AGTATTTGTTGAAATAGT 1182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 4552; 417pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAAAAAAAAAAAAAG
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ilarity 88.9%;
Conservative
                                                                                                                                                                                                                                      BP; 7 A; 2 C; 4 G; 7 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                               1.1%;
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Pred. No. 4.4e+02;
                                                                                                                                            Score 14.8;
Pred. No. 4
                                                                                                                         Miematches
                                                                                                                                                                                                                                      0 U; 0 Other;
                                                                                                                                                  4.4e+02;
                                                                                                                                                                           DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 20;
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                                                                                                                   Gaps
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RESULT

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ADK80450/c

ID ADK80450 standard; DNA; 20
XX
AC ADK80450;

XX
Chimeric phosphorothioate
XX
Navl.3; Analgesic; Nootrop
XX
Chimeric neuropathy; arthi
XX
Navl.3; Analgesic; Nootrop
XX
Infantile epilepsy; ataxi
OS
Synthetic.
XX
26-FEB-2004.
XX
PN
W02004016754-A2.
XX
PN
Roberds SL;
XX
PN
WPI; 2004-2003; 2003WO-US025-
XX
PN
Roberds SL;
XX
PN
Roberds SL;
XX
PN
WPI; 2004-203785/19.
XX
PN
Roberds SL;
XX
PN
Roberds SL;
XX
PN
Roberds SL;
XX
Chimayla, useful for useful
PT Wavl.3, e.g. pain, s
PT disorder, or ataxia.
XX
PS
Claim 4; SEQ ID NO 7784;
XX
PS
Claim 4; SEQ ID NO 7784;
XX
PS
Claim 4; SEQ ID NO 7784;
XX
Claim 4; SEQ ID NO 7784;
XX
PS
Claim 4; SEQ ID NO 7784;
XX
PS
Claim 4; SEQ ID NO 7784;
XX
Claim 4; SEQ ID NO 7784;
XX
PS
Claim 4; SEQ ID NO 7784;
XX
PD
CC compound and composition rel
CC sepcifically hybridizes w
CC compound and composition
CC associated with Navl.3, e.g. pain, s
CC neuropathic pain, post-he
CC pain from burns, migrain, c
CC pain from bu
      BXBXBXBXB
                                                                                                                                                                                                                                                                                          ADM69507
                                                                                                                                                                                                                                                                                                                       RESULT 795
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition associated with Nav1.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache, seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Nav1.3 expression, the oligonucleotides are designed to target different regions of the human Nav1.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache;
                                                               Plant gene polymorphism marker related primer, SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; SEQ ID NO 7784; 417pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chimeric phosphorothioate oligonucleotide to target Nav1.3
                                                                                                                                 03-JUN-2004
                                                                                                                                                                                                ADM69507;
                                                                                                                                                                                                                                                            ADM69507 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 10 A; 1 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-AUG-2002; 2002US-0403416P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               1252 TTTTGTTTTTTAATCAGA 1269
                                                                                                                                                                                                                                                                                                                                                                                                                           19 TTTTGATTTTTAATCACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16;
variation mapping; mutation mapping; plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                 (first
                                                                                                                                                                                                                                                            DNA;
                                                                                                                                 entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.8; DB 1;
Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         where the antisense compound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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RESULT 796
ADM69506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cc prepared from a plant homozygously having a mutation to be an object of the mapping; (b) A forward primer 1 containing a base corresponding to the gene polymorphic maker of one ecotype plant, a forward primer 2 containing a base corresponding to the genetic polymorphic maker of one ecotype plant, a forward primer 2 cc containing a base corresponding to the genetic polymorphism of the other ccotype plant and a reverse primer 3 based on the base sequence common cc with both the ecotype plants are prepared; (c) two kinds of coligonucleotides emitting fluorescence of different colors when the genetic polymorphism marker is detected are prepared; (d) an cc amplification reaction of the genomic DNA is carried out in the presence confidence of the primers 1, 2 and 3 and the two kinds of the oligonucleotides; (e) the fluorescence intensity emitted from the resultant reactional product is detected and (f) the position on the genome of the mutation site is continued from the results of detection. The present sequence is a cc primer, used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ঠ
                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a primer set and method for mapping least the variation/mutation part of a plant gene using a gene polymorphism marker. A mutation site of the plant gene is mapped by utilizing a genetic polymorphism marker as follows: (a) genomic DNA is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 7; SEQ ID NO 386; 120pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A primer set and method useful for mapping at least the variation/mutation part of a plant gene using a gene polymorphism marker.
               14-OCT-2003
                                                 JP2003289885-A
                                                                                                                Primer; variation mapping; mutation mapping; plant; gene polymorphism marker; ss.
                                                                                                                                                                  Plant gene polymorphism marker related primer, SEQ ID 385.
                                                                                                                                                                                                   03-JUN-2004
                                                                                                                                                                                                                                                                     ADM69506 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene polymorphism marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIKA ) RIKAGAKU KENKYUSHO
(SAIM-) SAI MEDIA KK.
(MATS/) MATSUI M.
                                                                                                                                                                                                                                                                                                                                                                                       1462 AGAAGGTGAACAATTTCA 1479
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                                                                                                                                                                                                                                                                                                                                                         18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.8; D
Pred. No. 4.4e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                         4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 20;
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RESULT 797
ADL33726/c
ID ADL3377
XX ADL337
XX ADL337
XX DT 03-JUN
XX LNA ol
XX Detect
XW Detect
XX Eynthe
XX Key
FT modifi
FT modifi
FT FT
FT T
XX X
PN W02004
XX W02004

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC utilizing a genetic polymorphism marker as follows: (a) genomic DNA is compared from a plant homozygously having a mutation to be an object of CC the gene polymorphic maker of one ecotype plant, a forward primer 2 containing a base corresponding to the gene polymorphic maker of one ecotype plant, a forward primer 2 containing a base corresponding to the genetic polymorphism of the other cectype plant and a reverse primer 3 based on the base sequence common CC with both the ecotype plants are prepared; (c) two kinds of CC oligonucleotides emitting fluorescence of different colors when the CC genetic polymorphism marker is detected are prepared; (d) an amplification reaction of the genomic DNA is carried out in the presence CC of the primers 1, 2 and 3 and the two kinds of the oligonucleotides; (e) CC the fluorescence intensity emitted from the resultant reactional product is detected and (f) the position on the genome of the mutation site is CC determined from the results of detection. The present sequence is a CC primer, used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                    LNA oligomer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       least the variation/mutation part of a plant gene using a gene by polymorphism marker. A mutation site of the plant gene is mapped by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 7; SEQ ID NO 385; 120pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A primer set and method useful for mapping at least the variation/mutation part of a plant gene using a gene polymorphism marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-FEB-2002; 2002JP-00025338
                    WO2004020575-A2
                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                  Detection; isolation; locked nucleic acid; LNA; ss
                                                                                                                                                                                                                                                                                                                                   03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                    ADL33726;
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(SAIM-) SAI MEDIA KK.
(MATS/) MATSUI M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention relates to a primer set and method for mapping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     l Similarity
16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AGAAGGTGGACACTTTCA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                        /*tag= b
/mod_base= OTHER
                                                                                                                                                                                                Location/Qualifiers
                                                 /mod_base= OTHER
/note= "Optionally biotinylated or 5′ AQ2-HEG3, where AQ
is anthraquinone and HEG is hexa-ethylene glycol"
                                                                                                                                    'note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; 3 C; 6 G; 4 T;
                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.1%;
88.9%;
                                                                                                                                  "Optionally
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8; D
Pred. No. 4.4e
0; Mismatches
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                                                                                                                                    LNA nucleotides'
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RESULT 798
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ID ADL596
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KW Gene t
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy; endothelial specific molecule-1; ESM-1; ESM-1 related disorder; diabetes; cancer; ischaemia; reperfusion injury; angiogenic disorder; immunological disorder; cardiovascular disorder; neurological disorder; antisense technology; ss.
                                                                                      modified_base
                                                                                                                                                                                             modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-JUN-2002; 2002US-0390928P
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                                                                                                                                                                                                                                /note= "OTHER= phosphorothioate backbone. All cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                               mod_base= OTHER
                                                                                                                                               mod_base=
                                                                                                                note= "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
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88.9%;
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        "OTHER= 2'-methoxyethyl (2'-MOE) nucleotides"
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Pred. No. 4.
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Best Local
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                                                                                                                                                                                                    cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotect gene therapy; endothelial specific molecule-1; ESM-1; ESM-1 related disorder; diabetes; cancer; ischaemia; reperfusion angiogenic disorder; immunological disorder; cardiovascular disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense compound, having a sequence targeted to a nucleic avencoding endothelial specific molecule-1 (ESM-1), useful for prepacemposition for treating e.g., diabetes, cancer or cardiovascular
                                                                                                              modified_base
                                                                                                                                                                                                                                                                            Human ESM-1 antisense oligonucleotide seqid 1952.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 15 A; 2 C; 1 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 3; SEQ ID NO 1935; 555pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disorder.
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                                         modified_base
                                                                                                                                                          Homo sapiens
                                                                                                                                                                                      neurological
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             /*tag=
/mod_ba
                                                     residues
                                                                                                                             Location/Qualifiers
                                                       /note= "OTHER= phosphorothioate residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                        entry)
              base= OTHER
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Pred. No. 4.4e
0; Mismatches
2'-methoxyethyl (2'-MOE)
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  nucleotides"
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                                                                       All cytidine
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KW ESM-1

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Best Local
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                                                                                                                                          cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; gene therapy; endothelial specific molecule-1; ESM-1; ESM-1 related disorder; diabetee; cancer; ischaemia; reperfusion injury; angiogenic disorder; immunological disorder; cardiovascular disorder; neurological disorder; antisense technology; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 16 A; 2 C; 1 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                        03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention describes a new antisense compound, having comprising 8-30 bp targeted to a nucleic acid encoding en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              encoding endothelial specific molecule-1 (ESM-1) composition for treating e.g., diabetes, cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2004021978-A2
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                                                                                                              sapiens
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/mod_ba
/mod_base= OTHER
/note= "OTHER= phosphorothioate
residues are 5-methylcytidines"
                                                                             Location/Qualifiers
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Pred. No. 4.
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                  backbone.
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endothelial
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                  cytidine
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RESULT 801
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention describes a new antisense compound, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), that specifically hybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition; inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury, or angiogenic, immunological, cardiovascular or neurological disorder. This sequence represents an antisense oligonucleotide that can be used to modulate expression of
                                                                                                        Human; NOVX; PCR; 88; congenital heart defect; cardiomyopathy; atherosclerosis; hypertension; pulmonary stenosis; scleroderma; adenocarcinoma; haemophilia; graft-versus-host disease; cancer; neurodegemerative disorder; Alzheimer's disease; Parkinson's di multiple sclerosis; diabetes; obesity; bronchial asthma; acquired immunodeficiency syndrome; AIDS; Crohn's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing composition for treating e.g., diabetes, cancer or cardiovascular
                                                                                                                                                                                                                                                                                                         Human NOVX PCR
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                                                                                      infectious disease;
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05-NOV-2001
06-NOV-2001
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21-NOV-2001
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29-NOV-2001
29-NOV-2002
16-MAY-2002
16-MAY-2002
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(ALSO/
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MILLER C E.
MISHRA V I.
MISHRA V I.
MISHRA U M.
PADIGARU M.
PADIGARU M.
PEYMAN J A.
REGER D K.
RIEGER D K.
ROTHENBERG M E.
SHENOV S G.
SHENOV S G.
SHIMKETS R A.
SPADERNA S K.
SPADERNA S K.
SPADERNA S K.
SPATRLING G G.
SPATRLING G G.
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BOLDOG F L.
BURGESS C E.
CASMAN S J.
CHAPOVAL A.
DHAWABAL M.
EDINGER S R.
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GROSSE W M.
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ETTENBERG S.
GANGOLLI E A.
GERLACH V.
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ALVAREZ E.
ANDERSON D W.
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MEZES P S.
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KHRAMTSOV N V.
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2001US-0335610P.
2001US-0333152P.
2001US-0333152P.
2001US-0333912P.
2001US-0334300P.
2002US-034807P.
2002US-0380968P.
2002US-0380968P.
2002US-0381043P.
2002US-0393148P.
2002US-0401479P.
2002US-0401479P.
2002US-0401626P.
2002US-04016393P.
2002US-0401693P.
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2001US-0333072P.
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RESULT 802
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XW micros
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                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
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              chimeric; antisense oligonucleotide; phosphorothioate; human; microseomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microseomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic, immunomodulator; cardiant; neuroprotective; antinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; immunomodulatory; cardiovascular; gene therapy; inflammation;
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sclerosis, di
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                                                                                                                                       Human mPGES-1 chimeric antisense oligonucleotide
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                                                                                                                                                                                                                                                                                  antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, carcer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                  Sequence 20
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16; Conser
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/note= "2'-O-methoxyethyls"
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/note= "2'-O-methocyethyls"
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                                                                                                                                            Score 14.8; DB 1; Length 20; Pred. No. 4.4e+02;
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RESULT 803

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ADM13994/c
The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mpCES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to CC 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1 in cells or tissues; and (3) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpCES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, cardiant, neuroprotective, cardiant, neuroprotective, vasotropic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or
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/mod base= OTHER
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/note= "phosphorothioate linkages and all cytidine
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                                                                               New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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residues are 5-methylcytidines"
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186; 132pp; English
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Pred. No. 4.4e+02
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e= "2′-O-methocyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                             base= OTHER
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                                                                                                                                                        "2'-O-methoxyethyls'
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   RESULT 806
ADM14002/c
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                                                                                                                                                 chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpcES-1; mpcES-1 inhibitor; microsomal prostaglandin E2 synthase; mpcES-1; mpcES-1 intidiabetic; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alibeimer's disease; arthritis; diabetes; cancer; ischammation; Alibeimer's disease; arthritis; diabetes; cancer; ischammation; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                Human mPGES-1
                                                                                                                                                                                                                                                                                                                                                                    01-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 4; SEQ ID NO 195; 132pp; English.
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                   modified_base
                                                                             Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                              chimeric antisense oligonucleotide SEQ
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Location/Qualifiers
1. .20
/*tag= b
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                                                                                                                                 neurological disorder;
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Pred. No. 4.4e
0; Mismatches
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RESULT 807
ADM14090/c
SEXSXEXE
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Best Local :
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              Human mPGES-1 chimeric antisense oligonucleotide
                                             01-JUL-2004
                                                                         ADM14090
                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4; SEQ ID NO 189; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ischemia.
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16; Conserv
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                                                                                                       standard; DNA; 20
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                                             (first entry)
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/mod_base= OTHER
/note= "2'-O-methoxyethyls"
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residues are 5-methylcytidines"
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e= "2'-O-methocyethyls"
                                                                                                                                                                                                                                                      88.9%;
                                                                                                                                                                                                                                                                       1.1%;
                                                                                                                                                                                                                                           <u>,</u>
                                                                                                                                                                                                                                                                       Score 14.8;
                                                                                                                                                                                                                                                           Pred.
                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                           No.
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The present sequence represents a chimeric antisense oligonucleotide carageted to human microsomal prostaglandin E2 synthase (mpCES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to Q43.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antisinflammatory, neuroprotective, nootropic, antisinflammatory, neuroprotective, nootropic, antistalianological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
Query Match
Best Local S
Matches 16
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                                                                                                                         Sequence
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residues are 5-methylcytidines"
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                            Score 14.8; DB 1
Pred. No. 4.4e+02
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                                                         DB 1; Length 20;
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ID ADMI41
AC AD
The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9434.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal
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                                                                                                                                                                                                                                                                                                                                                                               New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                      ID NO 338; 132pp; English.
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/note= "phosphorothioate linkages and
residues are 5-methylcytidines"
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e= "2'-O-methocyethyls'
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RESULT 809
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         WPI; 2004-305094/28.
                                                                                                                                                                                   25-SEP-2002; 2002US-0413549P
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                                                                                                                              (PHAA ) PHARMACIA CORP
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/note= "2'-O-methocyethyls"
16. .20
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e= "2'-O-methoxyethyls"
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Pred. No. 4.4e+02
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RESULT 810
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          THE FIRST SOLUTION AND ACCOUNT.
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Best Local
                                                                                                                                                                                                                                                                                                                                                            chimeric; antirellandin E2 synthase; mPGES-1; mrubo-1 .....ridiabetic; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; meuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; **mmunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chimeric; antisense oligonucleotide; phosphorothioate;
microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic ac
encoding mPGES-1, useful for preparing a composition for treating
inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                        modified_base
                                                                                                   modified_base
                                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                             cardiovascular
                                                                                                                                                                                                                                                                                                                                              immunomodulatory; cardiovascular; gene therapy; inflammation Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human mPGES-1 chimeric antisense oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADM14017 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 4; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADM14017;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ID NO 184; 132pp; English
                                                                                                                                                                                                                                                                                                                           disorder; neurological disorder; ss.
/note= "2'-O-methocyethyls"
16. .20
/*tag= c
                                                                                                                       /mod_base= OTHER
/note= "phosphorothioate linkages
residues are 5-methylcytidines"
                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.8; DB 1;
Pred. No. 4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                              and all cytidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NO:204
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RESULT 811
ADM14018/c
ID ADM140
XX ADM140
XX ADM140
XX ADM140
XX OLIMBER
CHIMBER
KW Chimer
KW Micros
KW Micros
KW Micros
KW Mimuno
KW Immuno
KW Immuno
KW Alzhei
KW reperf
KW cardic
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human mbGES-1 gene is located on chromosome 9, more specifically to 2934.3. The present invention also describes: (1) antisense compounds, chaving a sequence comprising 8-30 bp targeted to a nucleic acid encoding mbGES-1, which specifically hybridise with the nucleic acid mbGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mbGES-1 in cells or tissues; and (3) a method of treating an animal chaving a disease or condition associated with mbGES-1. Chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, antidiabetic, immunomodulatory and cardiovascular activities, and can be used as mbGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mbGES-1 e.g., inflammation, Alzhaimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
              chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpcES-1; mpcES-1 inhibitor; microsomal prostaglandin E2 synthase; mpcES-1; mpcES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                           Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                    01-JUL-2004
                                                                                                                                                                                                                                                                                                                                     ADM14018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; SEQ ID NO 204; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-SEP-2002; 2002US-0413549P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1520 AAAAAAAAAAAAGTAAAA 1537
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 Similarity
16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                                                                                                                                                                                                                     standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           to human microsomal prostaglandin E2 synthase (mPGES-1).
                                                                                                                                                                                                                                                                                                                                                                                                                                         AAAAAAAAAAAAAAA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence represents a chimeric antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               note= "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14.8; DB 1;
Pred. No. 4.4e+02;
    disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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CC targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The CC targeted to human mpGES-1 gene is located on chromosome 9, more specifically to CC 9434.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 by targeted to a nucleic acid encoding cmpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal CC antisense oligonuclectides and antisense compounds have cytostatic, cantisense oligonuclectides and antisense compounds have cytostatic, cantisense oligonuclective, noctropic, antiarthritic, vasotropic, antiinfiammatory, neuroprotective, noctropic, antiarthritic, vasotropic, cantinefiammatory, neuroprotective, noctropic, antiarthritics, and can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition associated with mpGES-1 e.g., inflammation, Alzheimer's contribute immunological, diabetes, cancer, ischaemia or reperfusion injury, or contributes ammunological, diabetes, cancer, ischaemia or reperfusion injury, or contributes immunological.
                                                                                                                                           Query Match
Best Local
                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.c. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
modified_base
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Synthetic.
                                                                                                                                                                                                                                   Sequence 20 BP; O A; O C; O G; 20 T; O U; O Other;
                                                                                                                                                                                                                                                                                                   ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 4; SEQ ID NO 205; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-305094/28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ischemia.
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                                                     1520 АЛАЛАЛАЛАЛАЛ 1537
20
                                                                                                                     16;
                                                                                                                                                 Similarity
AAAAAAAAAAAAAAA 3
                                                                                                                                                                                                                                                                                                   immunological, cardiovascular or neurological disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence represents a chimeric antisense oligonucleotide
                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pom/
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/note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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/note= "2'-O-methoxyethyls"
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                                                                                                                                                 88.9%;
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                                                                                                                                              Pred.
                                                                                                                                                                              Score
                                                                                                                     Mismatches
                                                                                                                                                    No.
                                                                                                                                                                              14.8;
                                                                                                                                                    .4e+02
                                                                                                                                                                                 DB 1;
                                                                                                                        0,
                                                                                                                        Gaps
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RESULT 812 ADM14088/c ID ADM140

ADM14088 standard; DNA;

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpCES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to gd34.3. The present invention also describes: (1) antisense compounds, mpGES-1, which specifically hybridise with the nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and composition is expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal can having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidabetic, immunomodulator, cardiant, neuroprotective, cardiant, neuroprotective, cardiant, neuroprotective, cardiant method search activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition associated with mpGES-1 e.g., inflammation, Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                               inflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-SEP-2002; 2002US-0413549P
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                                                                                                                                                                                                                                                                                                                             Claim 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:275
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                         2004-305094/28
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                                                                                                                                                                                                                                                                                                                               SEQ
                                                                                                                                                                                                                                                                                                                             ID NO 275; 132pp; English.
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/mod_base= OTHER
/note= "2'-O-methocyethyls"
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residues are 5-methylcytidines"
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3= "2'-O-methoxyethyls"
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RESULT 813
ADM14257/c
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Best Local :
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                      Claim 4; SEQ ID NO 444; 132pp; English
                                                           New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                    25-SEP-2003; 2003WO-US030374
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                                                                                                                                                                                                                                                                                                                                     modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:444
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                WPI; 2004-305094/28
                                                                                                                                                                  (PHAA ) PHARMACIA CORP
                                                                                                                                                                                          25-SEP-2002; 2002US-0413549P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chimeric; antisense oligonucleotide; phosphorothioate;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                                                                                                                                        note = "2' - 0 - methocyethyls"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                      /note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                 note= "2'-0-methoxyethyls"
                                                                                                                                                                                                                                                                                                                                                                  mod base= OTHER
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                                                                                                                                                                                                                                                                                                              _base= OTHER
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Pred. No. 4.
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The present sequence represents a chimeric antisense oligonucleotide

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RESULT 814
ADM14000/c
ID ADM140
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XX ADM140
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XX Onimer
KW Chimer
KW micros
KW m
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
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                                     25-SEP-2003; 2003WO-US030374.
                                                                                                   08-APR-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disorder; neurological disorder; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
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e= "2'-O-methoxyethyls"
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Pred. No. 4.4e+02;
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                    chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:193
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                                                                                                              modified_base
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Pred. No. 4.4e+02;
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/note= "phosphorothioate linkages and all cytidine

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ADM14014/c
ID ADM140
XX
AC ADM140
DT 01-JUL
XX
DB Human
XX
KW chimer
KW micros
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                                                                                                                                                                                                                                                                                                                                                         RESULT 816
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                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
chimeric; antisense oligonucleotide; phosphorothioate; human;
microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor;
                                                Human mPGES-1
                                                                                                                                                ADM14014 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                         Sequence
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                                                chimeric antisense oligonucleotide SEQ ID
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/note= "2'-O-methocyethyls"
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/note= "2'-O-methoxyethyls"
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Pred. No. 4.4e+02;
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1520 AAAAAAAAAAAAGTAAAA 1537

Matches Query Match Best Local :

16;

Conservative

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4.4e+02 DB 1;

Length

20

Indels

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Gaps

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Local

Similarity

88.9%;

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The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The Chuman mpGES-1 gene is located on chromosome 9, more specifically to CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1 in cells or tissues; and (3) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC mpGES-1 in cells or condition associated with mpGES-1. whose simple condition associated with mpGES-1. chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, recomposition associated with mpGES-1 antisense compound can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC chitchia artiritis, diabetes, cancer, isbaemia or reperfusion injury, or condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition associated with mpGES-1 e.g., inflammation, Alzheimer's conditions artiritis, diabetes, cancer, isbaemia or reperfusion injury, or conditions and conditions and conditions are composition or reperfusion injury, or conditions are composition or conditions or reperfusion injury, or conditions are composition or conditions or reperfusions injury, or conditions are composition or conditions or conditions are compositions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
    Sequence 20
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0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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residues are 5-methylcytidines"
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                         targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric
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                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.cinflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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/note= "phosphorothicate linkages and all cytidine
/residues are 5-methylcytidines"
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antisense oligonucleotides and antisense compounds

have cytostatic,

New antisense compound,

having a sequence targeted

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nucleic

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RESULT 818
ADM13991/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1 mpGES-1 inhibitor; microsomal prostaglandin E2 synthase; mbjas-1; mctor; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                           Key
modified_base
                           WPI; 2004-305094/28.
                                                     Gierse JK;
                                                                                                                                                                                                                                                                                                                        modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:178.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADM13991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                             25-SEP-2002; 2002US-0413549P
                                                                                                                                       25-SEP-2003; 2003WO-US030374
                                                                                                                                                                    08-APR-2004.
                                                                                                                                                                                               WO2004028458-A2
                                                                                                                                                                                                                                                                modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cardiovascular
                                                                                 (PHAA ) PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 AAAAAAAAAAAAAAAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                   /note=
16. .2
                                                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "phosphorothioate linkages and
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                     *tag=
                                                                                                                                                                                                                                                                                               mod_
                                                                                                                                                                                                                                                                                                            /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
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                                                                                                                                                                                                                         base= OTHER
== "2'-O-methoxyethyls"
                                                                                                                                                                                                                                                                                               base= OTHER
                                                                                                                                                                                                                                                                                "2'-O-methocyethyls"
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ARESULT 819
ADM1400
XX
ADM140
AC ADM140
XX
ADM140
XX
ADM140
XX
Chimer
KW micros
KW micros
KW immuno
KW immuno
KW reperf
KW cardio
XX
Chimer
KW reperf
KW rep
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC 9344.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 by targeted to a nucleic acid encoding CC mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mPGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mPGES-1. MPGES-1 chimeric CC antisense oligonuclectides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, antiarthritic, vasotropic, CC ophthalmological, immunomodulatory and cardiovascular activities, and can be used for preparing a composition for treating a disease or CC condition associated with mPGES-1 entisense compound CC can be used for preparing a composition for treating a disease or CC condition associated with mPGES-1 e.g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzhelmer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:190.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADM14003 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cargeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) APPLICATION AND AND AND ASSESSED.
                                                                                      modified_base
                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                        modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                             /mod_base= OTHER
/note= "2'-O-methocyethyls"
16. 20
                                                                                                                                                                                                                                          /mod_base= OTHER
/note= "phosphorothioate linkages and
reaidues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
/mod_base= OTHER
/note= "2'-O-methoxyethyls"
                                                             /*tag=
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Pred. No. 4.4e
O; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                 all cytidine
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RESULT 820
ADM14005/c
ID ADM140
XX
AC ADM140
XX
AC ADM140
XX
DT 01-UUL
XX
Human
XX
Chimer
KW micros
KW micros
KW neurop
KW immuno

Human mPGES-1 01-JUL-2004

chimeric antisense oligonucleotide SEQ ID

NO:192

antidiabetic;

(first entry)

ADM14005 standard; DNA; 20

ADM14005

chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpcBS-1; mpGBS-1 inhibitor; microsomal prostaglandin E2 synthase; mpcBS-1; mcGBS-1 inhibitor; cardiant E2 synthase inhibitor; cytostatic; antidiabetic immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;

cardiovascular

disorder; neurological disorder;

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20 AAAAAAAAAAAAAAAAAAA

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                                                                                                                                   The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpdES-1). The CC human mpGES-1 gene is located on chromosome, by more specifically to Q934.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid encoding cd inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cyrostatic, cophthalmological, immunomodulator, cardiant, neuroprotective, notropic, ophthalmological, immunomodulator, and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                         Query Match
Best Local
                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.dinflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                 Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; SEQ ID NO 190; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-SEP-2002; 2002US-0413549P
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1520 AAAAAAAAAAAGTAAAA 1537
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                                                     16;
                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PHARMACIA CORP
                                                     Conservative
                                                                       88.9%;
                                                                                          1.18;
                                                     0; Mismatches
                                                                                          Score 14.8;
                                                                         Pred. No. 4.4e+02
                                                                                              DB 1;
                                                                                          Length
                                                     0
                                                       Gaps
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RESULT 821
ADM14246/c
ID ADM142
XX
AC ADM142
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                                                                                                                                                                             Matches
                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
   ADM14246;
                                  ADM14246 standard; DNA; 20
                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; SEQ ID NO 192; 132pp; English.
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                                                                                                                                                                                              Local Similarity
                                                                                                                                               1520
                                                                                                                  20
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                                                                                                                                                                                                                                            20
                                                                                                                  AAAAAAAAAAAAAAAAA 3
                                                                                                                                                                                                                                            BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                               Conservative
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Pred. No. 4.4e+02
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antidiabetic, immunomodulator, cardiant, neuroprotective, vasotropic, antiarthritic, vasotropic, antiarthritic, vasotropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's condition associated with mPGES-1 e.g., inflammation, alzheimer's mediasese, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                    The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues, and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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ophthalmic,

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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modified_base
                                                                                                                                                                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
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                                                                                                                                                                                                                        WO2004028458-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADM14011;
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16; Conserv
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/mod_base= OTHER
/note= "phosphorothioate linkages
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                            /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag= a
/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                      base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                "2'-O-methocyethyls"
                                                                                                                                                                                                                                                                                     "2'-O-methoxyethyls'
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0; Mismatches
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    antisense compounds,

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25-SEP-2002; 2002US-0413549P 25-SEP-2003; 2003WO-US030374.

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RESULT 824
ADM14270/c
ID ADM142
XX ADM142
XX ADM142
XX O1-JUL
XX Human
XX Chimex
KW micros
KW mi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents a chimeric antisense oligonucleotide cargeted to human microsomal prostaglandin E2 synthase (mpGES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to CC 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding cmpGES-1, which specifically hybridise with the nucleic acid encoding cmpGES-1 in cells or tissues; and (3) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal chaving a disease or condition associated with mpGES-1. MpGES-1 chimeric antidiabetic, immunomodulator, cardiant, neuroprotective, cantidiabetic, immunomodulator, cardiant, neuroprotective, antiarithical, neuroprotective, nootropic, antiarithical, immunomodulatory and cardiovascular activities, and can be used as mpCES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating Alzheimer's composition associated with mpGES-1 e.g., inflammation, Alzheimer's composition for treating a cardiary, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chimeric; antisense oligonucleotide; phosphorothioate; human; microscmal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microscmal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological;
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                                                                                                                                                                                                                                                                                                                                                                                                                              immunomodulatory; cardiovascular; gene therapy; inflammation;
Alzheimer's disease; arthritis; diabetes; cancer; ischaemia;
reperfusion injury; ophthalmic disorder; immunological disorder;
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       modified_base
                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                          cardiovascular
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chimeric antisense oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                             disorder; neurological disorder; ss
                                              residues
                                                                                                                                                                                                                    Location/Qualifiers
                                           /note= "phosphorothioate linkages and
residues are 5-methylcytidines"
                                                                                                               /*tag= b
/mod_base= OTHER
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RESULT 825
ADM14009/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local :
chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antid immunomodulator; cardiant; neuroprotective; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compound, having a sequence targeted to a nucleic avencoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                   Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:196
                                                                                                                                                                                                              ADM14009;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; SEQ ID NO 427; 132pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                           standard;
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16. .20
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/mod_base= OTHER
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/mod_base= OTHER
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                                                                                                                                                                entry)
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                                                                                                                                                                                                                                                           ВP
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Pred. No. 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.4e+02
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                          antidiabetic;
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CC The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The CC human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC maying a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, nouroprotective, cc antidiabetic, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense or CC condition associated with mpGES-1 engaged a disease or CC condition associated with mpGES-1 engaged and cardiovascular activities, and can be used for preparing a composition for treating a disease or CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                          Matches
                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic au encoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; SEQ ID NO 196; 132pp; English.
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                                                                                                                                                                                                                     Sequence
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                                                                                                                                    Similarity
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                                                   AAAAAAAAAAAGTAAAA 1537
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                                                                                                          Conservative
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/note= "2'-O-methocyethyls"
16. .20
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                                                                                                       Score 14.8; D
Pred. No. 4.4e
O; Mismatches
                                                                                                                                                                                                                     T; 0 U;
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mBGES-1 gene is located on chromosome 9, more specifically to 9334.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mBGES-1, which specifically hybridise with the nucleic acid mBGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mBGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mBGES-1. MBGES-1 chimeric
                                                                                                                                                                                                                                            New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.g. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
                         antisense oligonucleotides and antisense compounds have cytostatic,
                                                                                                                                                                                                 Claim 4; SEQ ID NO 197; 132pp; English.
                                                                                                                                                                                                                                                                                                                                          Gierse JK
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                   immunomodulator,
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    neuroprotective,
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                   cardiant, neuroprotective,
    antiarthritic,
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   vasotropic
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RESULT 827
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
   New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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Pred. No. 4.4e+02;
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Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                             modified_base
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/mod_base= OTHER
/note= "2'-O-methocyethyls"
                 /*tag= c
/mod_base= OTHER
/note= "2'-O-methoxyethyls"
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/note= "phosphorothicate linkages and all cytidine
residues are 5-methylcytidines"
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Pred. No. 4.4e+02;
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WO2004028458-A2.

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ADM14075/C
ID ADM140
XX ADM140
XX ADM140
XX ADM140
XX ADM160
DT 01-JUL
XX Chimer
XX Chimer
XX Immuno
XX micros
XW micros
XW micros
XW micros
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Best Local &
                                                                                                     chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzhelmer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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                                                      Homo sapiens
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RESULT 830
ADM14189/c
ID ADM141
XX
AC ADM141
XX
AC ADM141
XX
OT 01-JUL

standard; DNA;

20 ВP

ADM14189; ADM14189

01-JUL-2004

(first entry)

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                                                                                                                                  The present sequence represents a chimeric antisense oligonucleotide CC targeted to human microsomal prostaglandin E2 synthase (mpCES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to compare the present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid encoding CC mpGES-1 in cells or tissues; and (3) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC mpGES-1 in cells or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective, cardiant, neuroprotective, or antidiabetic, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's condition associated with mpGES-1 e.g., inflammation, Alzheimer's contributed artificial dispater.
                                                                                Query Match
Best Local
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                                                                                                                                  Sequence 20
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                                                                  Conservative
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residues are 5-methylcytidines"
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Human mPGES-1 chimeric antisense oligonucleotide

SEQ ID

NO:376

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antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                           The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic,
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l prostaglandin E2 synthase; mPGES-1; mPGES-1
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Matches 16
The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encodi
                                                                                                                                                                Claim
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microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor;
microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic;
                                                                                                                                                                                                                                              New antisense compound, having a sequence targeted to a nucleic acencoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                               Gierse
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                                                                                                                                                                   4; SEQ ID NO 183;
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Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
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llarity 88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC human mpGES-1 gene is located on chromosome 9, more specifically to G234.3. The present invention also describes: (1) antisense compounds, CC having a sequence comprising 8-30 by targeted to a nucleic acid encoding CC mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and CC inhibits its expression; (2) a method of inhibiting the expression of CC mpGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mpGES-1. MpGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cytostatic, CC antidiabetic, immunomodulator, cardiant, neuroprotective, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or CC condition associated with mpGES-1 e.g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:191
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Pred. No. 4.4e+02;
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                chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpcBS-1; mpcBS-1 inhibitor; microsomal prostaglandin E2 synthase; mpcBS-1; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; antiinflammatory; neuroprotective; antiarthritic; vasotropic; ophthalmological;
                                                                                                                                                           01-JUL-2004
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                                                                                                                                                           (first entry)
                                                                                                                         chimeric
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Query Match
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Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9344.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid meGES-1 and inhibits its expression, (2) a method of inhibiting the expression of mPGES-1 in cells or tissues, and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric
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                                                                                                                                                                                                                 Sequence 20
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                                                                                                                                                                                                                                                                       ophthalmic,
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Pred. No. 4.4e+02;
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                                                                                                                         The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective.
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                                              antiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and activities, acti
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                              residues are
                                                                                                                                                                                                               /notē= "phosphorothioate linkages and all cytidine
residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                       /mod_base= OTHER
/note= "2'.-O-methoxyethyls"
                                                                                                        /*tag=
                                                                                                                                                        mod_base= OTHER
                                                                                                                                                                                                                                                  mod_base= OTHER
                                                                                                                                         note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nce represents a chimeric antisense oligonucleotide microsomal prostaglandin E2 synthase (mPGES-1). Th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.1%;
                                                                                                                                           "2'-O-methocyethyls"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14.8;
Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 20;
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ADM14388/c
ID ADM143
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The CC human mPGES-1 gene is located on chromosome 9, more specifically to Q934.3. The present invention also describes: (1) antisense compounds, CC mPGES-1, which specifically hybridise with the nucleic acid encoding CC inhibits its expression; (2) a method of inhibiting the expression of CC mPGES-1 in cells or tissues; and (3) a method of treating an animal CC having a disease or condition associated with mPGES-1. MPGES-1 chimeric CC antisense oligonucleotides and antisense compounds have cyrostatic, cophthalmological, immunomodulator, cardiant, neuroprotective, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound CC can be used for preparing a composition for treating a disease or CC condition associated with mPGES-1 e-g., inflammation, Alzheimer's CC disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 16
                                                                                                      chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compound, having a sequence targeted to a nucleic avenceding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                     Synthetic
                                                                                                                                                                                                                                                                                               Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:575
                                                                                                                                                                                                                                                                                                                                     01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                          ADM14388;
                                                                                                                                                                                                                                                                                                                                                                                                            ADM14388 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-SEP-2002; 2002US-0413549P
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                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1520 АААААААААААСТААА 1537
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4; SEQ ID NO 208; 132pp; English.
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modified_base

Location/Qualifiers
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RESULT 839
ADM14013/c
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AC ADM140
AC ADM140
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DT 01-JUL
XX
DT Human
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                                                                                                                                                                                                                                      Query Match
Best Local :
                                                                                                                                                                                                                        Matches
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 Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:200.
                              01-JUL-2004
                                                                                                                                                                                                                                                                                 Sequence
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                                                           ADM14013;
                                                                                         ADM14013 standard;
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                                                                                                                                                                                         1520 AAAAAAAAAAAAGTAAAA 1537
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                                                                                                                                                                                                                        Conservative
                             (first entry)
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/note= "2'-O-methocyethyls"
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= "2'-O-methoxyethyls"
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                                                                                                                                                                                                                                    Score 14.8; DB 1; Length 20; Pred. No. 4.4e+02;
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to g934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid encoding conhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal chaving a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisinse oligonucleotides and antisense compounds have cytostatic, and cantiinflammatory, neuroprotective, nootropic, antiarthritic, vasotropic, antisinflammatory, neuroprotective, nootropic, antistatinglacial, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mpGES-1 e.g., inflammation, Alzheimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or ophthalmic, immunological, cardiovascular or neurological disorder.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense compound, having a sequence targeted to a nucleic acencoding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 4; SEQ ID NO 200; 132pp; English.
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/note= "2'-O-methoxyethyls"
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Query Match Best Local Similarity

1.1%;

Score 14.8; DB 1; Pred. No. 4.4e+02;

DB 1;

Length

20;

Sequence 20

BP;

0 A; 0 C; 0 G;

20 T; 0 U; 0 Other;

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RESULT 840
ADM14019/c
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         The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostaglandin E2 synthase (mPGES-1). The human mPGES-1 gene is located on chromosome 9, more specifically to 9q34.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and
                                                                                                                                                New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e.ginflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                         Claim 4; SEQ ID NO 206; 132pp; English.
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expression:
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residues are 5-methylcytidines"
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RESULT 841
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Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense origonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation;
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                                                                                    (PHAA ) PHARMACIA CORP
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residues are 5-methylcytidines"
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Pred. No. 4.4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                            chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mPGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase; inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder; cardiovascular disorder; neurological disorder; ss.
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                                                                                                                                                                                                                                                                                                                                  Synthetic.
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Pred. No. 4.4e+02;
0; Mismatches 2
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RESULT 843
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chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1 inhibitor; microsomal prostaglandin E2 synthase; mpGES-1; mpGES-1; mtdiabetic; antidiabetic; cardiant; neuroprotective; antiinflammatory; neuroprotective; notropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
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                                                                                                                                                                                                                                      Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:180
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                                                                                                                                                                                                                                                                                                                                                                                             standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88.9%;
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Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
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RESULT 844
ADM13998/c
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                                                                                                                            Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                human mPGES-1 gene is located on chromosome 9, more specifically to 9934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding mPGES-1, which specifically hybridise with the nucleic acid mPGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mPGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mPGES-1. MPGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antidiabetic, immunomodulator, cardiant, neuroprotective,
                                                                                                                                                                                                                                     antidiabetic, immunomodulator, cardiant, neuroprotective, antiinflammatory, neuroprotective, mootropic, antiarthritic, vasotropic, ophthalmological, immunomodulatory and cardiovascular activities, and can be used as mPGES-1 inhibitors and in gene therapy. The antisense compound can be used for preparing a composition for treating a disease or condition associated with mPGES-1 e.g., inflammation, Alzhaimer's disease, arthritis, diabetes, cancer, ischaemia or reperfusion injury, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                        ophthalmic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; SEQ ID NO 180; 132pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  targeted
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                                                                                                                                            Similarity
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                                                                                            ААААААААААСТААА 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             to human microsomal prostaglandin E2 synthase (mPGES-1).
                                                                AAAAAAAAAAAAAAAAA 3
                                                                                                                             Conservative
                                                                                                                                                                                          BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence represents a chimeric antisense oligonucleotide
                                                                                                                                                                                                                        immunological,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disorder; neurological disorder; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "phosphorothicate linkages residues are 5-methylcytidines"
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                                                                                                                                         1.1%;
88.9%;
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                                                                                                                                                                                                                          cardiovascular
                                                                                                                             0
                                                                                                                                            Score 14.8; DB 1;
Pred. No. 4.4e+02;
                                                                                                                          Mismatches
                                                                                                                                                             DB 1;
                                                                                                                                                                                                                      or neurological
                                                                                                                                                           Length 20;
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                                                                                                                             Gaps
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The present sequence represents a chimeric antisense oligonucleotide targeted to human microsomal prostagiandin EZ synthase (mpGES-1). The human mpGES-1 gene is located on chromosome 9, more specifically to 934.3. The present invention also describes: (1) antisense compounds, having a sequence comprising 8-30 by targeted to a nucleic acid encoding mpGES-1, which specifically hybridise with the nucleic acid mpGES-1 and inhibits its expression; (2) a method of inhibiting the expression of mpGES-1 in cells or tissues; and (3) a method of treating an animal having a disease or condition associated with mpGES-1. MpGES-1 chimeric antisense oligonucleotides and antisense compounds have cytostatic, antisense oligonucleotides and antisense compounds have cytostatic, antisinflammatory, neuroprotective, nootropic, antistrifical, immunomodulatory and cardiovascular activities, and can be used as mpGES-1 inhibitors and in gene therapy. The antisense compound cardiovascular activities, and can be used for preparing a composition for treating a disease or
                                                                                                                                                                                                                                                                                                                                   New antisense compound, having a sequence targeted to a nucleic adenooding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   microsomal prostaglandin E2 synthase; mpGES-1; mPGES-1 inhibitor; microsomal prostaglandin E2 synthase inhibitor; cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; antinflammatory; neuroprotective; antinflammatory; neuroprotective; notropic; ophthalmological;
                                                                                                                                                                                                                                                                                      Claim 4; SEQ ID NO 185;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Gierse JK;
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                                                                                                                                                                                                                                                                                                                         1schemia.
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 be used for preparing dition associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense oligonucleotide; phosphorothioate;
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/mod_ba
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residues are 5-methylcytidines"
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== "2'-O-methocyethyls"
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                                                                                                                                                                                                                                                                                    132pp; English.
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mPGES-1 e.g.,
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RESULT 845
ADM14007/c
ID ADM140
XX ADM140
XX ADM140
XX Chimer
KW chimer
KW micros
KW m
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chimeric; antisense oligonucleotide; phosphorothioate; human; microsomal prostaglandin E2 synthase; mpCES-1; mpCES-1 inhibitor; microsomal prostaglandin E2 synthase; mpCES-1; mpCES-1; mtidiabetic; immunomodulator; cardiant; neuroprotective; antiinflammatory; neuroprotective; nootropic; antiarthritic; vasotropic; ophthalmological; immunomodulatory; cardiovascular; gene therapy; inflammation; Alzheimer's disease; arthritis; diabetes; cancer; ischaemia; reperfusion injury; ophthalmic disorder; immunological disorder;
                                                                                                                                                                                                                                                                                                                                                                                                            25-SEP-2003; 2003WO-US030374.
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                                                                                                  New antisense compound, having a sequence targeted to a nucleic acid encoding mPGES-1, useful for preparing a composition for treating e. inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
                                                                                                                                                                                                                                                                                                                                                            25-SEP-2002; 2002US-0413549P
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                          Claim 4; SEQ
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residues are 5-methylcytidines"
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                          194; 132pp; English
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Pred. No. 4.
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RESULT 846
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CHIMER
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Best Local :
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Synthetic.
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25-SEP-2003; 2003WO-US030374.

08-APR-2004.

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                                                                         modified_base
                                                                                                                                                                                                  Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human mPGES-1 chimeric antisense oligonucleotide SEQ ID NO:403
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Pred. No. 4.4e+02;
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Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense compound, having a sequence targeted to a nucleic avenceding mPGES-1, useful for preparing a composition for treating inflammation, Alzheimer's disease, arthritis, diabetes, cancer or
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                             Human oligonucleotide #1790.
                                                            15-JUL-2004
                                                                                            ADO46424;
                                                                                                                         ADO46424 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                  Sequence 20
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                                                                                                                                                                                                          20 AAAAAAAAAAAAAAAA 3
                                                                                                                                                                                                                                                                     16;
                                                                                                                                                                                                                                                                                    Similarity
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16. .20
/*tag= c
/mod_base= OTHER
/~~te= "2'-O-methoxyethyls"
                                                           (first entry)
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/mod_base= OTHER
/note= "2'-O-methocyethyls"
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residues are 5-methylcytidines"
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Pred. No. 4.
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Human; ss;

interleukin-4 receptor; IL-4; interleukin-5 receptor;

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chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)

-5 receptor, CCR1, CCR3, Botaxin-1, RANTES, MCP4, CD23, ICAM, VCAM,

C tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention

C also relates to a method of screening a candidate compound that binds to

come or more nucleic acid target(s) or expressed product(s), for the

prevention and/or treatment of a respiratory or lung disease. The

c oligonucleotides are useful for reducing or inhibiting expression of a

gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor,

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a,

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a,

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a,

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a,

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a.

CCR1, CCR3, Extaxin-1, RANTES, MCP4, CD23, ICAM, CR2, CD23, ICAM, CR2, CD23, ICAM, CR2, CD23, ICAM, CR2, CD23, ICAM, CD23, IC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                        allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shahabuddin
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23-APR-2002;
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     bronchoconstriction.
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(SAND/)
(TANG/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AGUILAR D.
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in S, Lu H,
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2002WO-US013143.
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This sequence represents an oligonucleotide of the
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Cong H;
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1520 AAAAAAAAAAAAGTAAAA 1537

Best Local Similarity Matches 16; Conserv

Conservative

0;

Mismatches

1.1%; 86.9%;

Score 14.8; DB 1; Pred. No. 4.4e+02;

DB 1;

Length Indels

20;

0,

0,

Query Match

Sequence 20

BP; 0 A; 1 C; 0 G; 19 T; 0 U; 0 Other;

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RESULT 849
ADO03711
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                                                                                                                                                          The invention describes a reagent (1) comprising a particle bound to at Cleast one Raman label and a specific binding member, where the Raman CC label can be activated to provide a surface-enhanced Raman scattering CC (SERS) effect or comprising a specific binding member having two or more CC different Raman labels bound to it. Also described are: a test kit (II), CC comprising (I) in one container and a silver, gold or copper Raman CC enhancer stain in another container, and a fibre optic detection device (III), having a bundle of optical fibres terminating with ends of the CC pitical fibre, where a several of the optical fibres have (I) located at CC the ends of the optical fibre. (I) is useful for: detecting for the CC presence or absence of one or more target analytes in a sample, the CC target analytes having at least two binding sites; detecting the presence CC or absence of one or more target nucleic acid in a sample, the sequence of the nucleic acid having at least two portions; and for screening one CC or more molecules to determine whether the molecule is a ligand to one or more specific receptors. This sequence represents an oligonucleotide associated with the SERS-based detection analyte detection method.
                                                                 Matches
                                                                               Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reagent, useful for detecting target analyte e.g., nucleic acid, comprising particle having bound to at least one Raman label, which activated to provide surface-enhanced Raman scattering effect, specific binding member.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-MAY-2002; 2002US-0378538P
28-MAY-2002; 2002US-0383630P
14-JUN-2002; 2002US-00172428
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SERS-based analyte detection oligonucleotide segid 31.
                                                                                                                                Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; SEQ ID NO 31; 55pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-418413/39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MIRK/) MIRKIN C
(CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88
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1 AAAAAAAAAAAAAAAAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cao Y,
                                                                                                                                вP;
                                                                 Conservative
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Pred. No. 4.4e+02;
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RESULT 850
ADP78301/c
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RESULT 851
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a compound which specifically hybridizes with a nucleic acid molecule encoding GFAT, and inhibits the expression of GFAT. Specifically claimed are antisense oligonucleotides capable of modulating the expression of GFAT, and which comprise any of the 3063 sequences of 20 base pairs, given in the specification. The compound, composition and methods are useful for treating a disease or condition associated with GFAT, such as a disease or condition, e.g. diabetes, a cardiovascular or neurological disorder, ischemia/reperfusion injury. They are also useful in research and diagnostics for modulating the expression of GFAT. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2'-MOE wings and a deoxy gap, these oligonucleotides inhibit human GFAT expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding glutamine-fructose-6-phosphate amidotransferase (GFAT), for treating diabetes, a cardiovascular or neurologic disorder. ischemia/reperfusion injury.
                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                TTTTGTTTTTTAATCAGA 1269
                                                                                                                                                                                                                                                                                                                                                                20 BP; 11
                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Crosby SD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "
17. .20
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/mod_base= other
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mod_base= other
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                                                                                                                                                                                                                                                                                                                                                             A; 1 C; 2 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                      88.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "2-methoxyethyl wing"
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                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                      Score 14.8; DB 1
Pred. No. 4.4e+02
                                                                                                                                                                                                                                                                                                  DB 1; Length 20;
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RESULT 852
ADP20152
ID ADP201
XX
AC ADP201
XX
DT 26-AUG
XX
DE Nuclei
XX
Nuclei
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                      rejection, e.g. cardiac or kidney transplant rejection, in an individual comprises detecting the expression level of one or more genes. The methods, system and kits are useful in diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in a xenotransplant rejection or mechanical organ replacement rejection, in a individual. The method is also useful in assessing the immune status of an individual. The methods are also useful in diagnosing and monitoring diseases that involve the immune system, e.g. rheumatoid arthritis, lupus, inflammatory bowel diseases, multiple sclerosis, HIV/AIDS or viral, bacterial or fungal infection. The present sequence represents a primer for a 50 mer oligonucleotide marker for diagnosis and monitoring of allograft rejection and other disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to diagnosing or monitoring transplant rejection, e.g. cardiac or kidney transplant rejection, in an individual comprises detecting the expression level of one or more genes. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an individual, comprises detecting the expression level of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transplant rejection; immune system; rheumatoid arthritis; lupus; inflammatory bowel disease; multiple sclerosis; HIV; AIDS; ss; primer.
                                 Nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 58; SEQ ID NO 755; 1762pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-400724/37.
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20-DEC-2002; 2002US-00325899
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                                                                 26-AUG-2004
                                                                                                  ADP20152;
                                                                                                                                ADP20152 standard;
                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 0 A; 3 C; 8 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rosenberg
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                                                                                                                                                                                                                                                                                16;
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                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                Conservative
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                                 detection method linking oligonucleotide #66.
                                                                 (first
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                                                                                                                                  DNA;
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                                                                entry)
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                                                                                                                                  ВP
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                                                                                                                                                                                                                                                                                               Score 14.8; DB 1
Pred. No. 4.4e+02
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                                                                                                                                                                                                                                                                                 Gaps
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Nucleic acid

detection;

nanoparticle-oligonucleotide conjugate;

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RESULT 853
ADP20137
ID ADP201
XX
AC ADP201
XX
DT 26-AUG
DT 26-AUG
XX
CX
DE Nuclei
XX
Nuclei
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                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method of detecting a nucleic acid with at CC least two portions by providing a type of nanoparticle-oligonucleotide CC conjugate, contacting the nucleic acid and nanoparticles to allow CC hybridisation of the oligonucleotides with the two or more portions of CC the nucleic acid and observing a detectable change brought about by Chybridisation. The oligonucleotides have a sequence complementary to the Sequence of at least two portions of the nucleic acid. Hybridisation of CC in a detectable change. The method is used for detection and separation CC in a detectable change. The method is used for detection and separation CC in a detectable change. The method is used for detection and separation CC in uccleic acids (e.g. viral DNA, synthetic DNA, structurally-modified DNA, DNA CC from biological sources or PCR products) for diagnosis of various CC diseases (such as genetic diseases, bacterial infections and viral CC monitoring gene therapy. This sequencing, paternity testing and CC coligonucleotide of the invention.
                                                                                                                                                                                                                                                                                     Query Match
Best Local
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13-JAN-2000;
26-APR-2000;
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 Nucleic acid detection; nanoparticle-oligonucleotide conjugate;
                                Nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-JUN-2004
                                                                                                 ADP20137;
                                                                                                                             ADP20137 standard; DNA;
                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U;
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DNA sequencing; paternity testing; linking oligonucleotide; ss.
                                                                 26-AUG-2004
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26-JUN-2000;
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29-JAN-1999;
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                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                        AAAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                            AAAAAAAAAAAAA 18
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99US-0034467.
2000US-0176499P.
2000US-0200161P.
2000US-0603830.
2000US-0213966P.
2001US-00760500.
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                                                                                                                                                                                                                                                                       Conservative
                               detection method linking oligonucleotide #54.
                                                                (first entry)
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                                                                                                                                                                                                                                                                    Score 14.8; DB 1; Length 20; Pred. No. 4.4e+02; O; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                     0 Other;
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                                                                                                                                                                                                                                                                       Gaps
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SXBXBXBXB

09-SEP-2004 ADP69379

(first entry)

ADP69379 standard; DNA;

20

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human; antisense oligonucleotide; mitochondrial membrane; Human mitoNEET-specific antisense oligonucleotide #273.

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RESULT 854
ADP69379/c
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                                                                                                                                  Matches
                                                                                                                                                               Query Match
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29-JUN-1999;
25-JUN-1999;
13-JAN-2000;
26-APR-2000;
26-JUN-2000;
26-JUN-2000;
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                                                                                                                                                                                                                                oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nanoparticles useful for detection and separation of nucleic acids e.g. genes associated with disease, in a diagnostic assay, comprise several oligonucleotides attached to them.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   genetic disease; bacterial infection; viral infection; forensic; DNA sequencing; paternity testing; linking oligonucleotide; ss.
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                                                                                               1520 AAAAAAAAAAAAGTAAAA 1537
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                                                                                                                                                 Similarity
                                                                  AAAAAAAAAAAAAAA 18
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Garimella V, Li Z;
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; 2000US-0200161P.
; 2000US-00603830.
; 2000US-0213906P.
; 2001US-00760500.
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                                                                                                                                  Conservative
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99US-00344667.
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                                                                                                                                              1.1%;
88.9%;
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                                                                                                                                0,
                                                                                                                                               Score 14.8; DB 1; Length Pred. No. 4.4e+02;
                                                                                                                                  Mismatches
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                                                                                                                                              4.4e+02;
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                                                                                                                                                                                                 0 Other;
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                                                                                                                                  Indels
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RESULT 855
ADF69506/c
ID ADF695
XX ADF695
XX ADF695
XX O9-SEP
XX Human;
XW human;
XW insuli
XW inmuno
XW neurol
XW remet
XX OS Homo s
XX Homo s
XX PD 24-JUN
XX
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Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        e.g.
                                                                                                                      human; antisense oligonucleotide; mitochondrial membrane; diabetes; insulin sensitising antidiabetic thiazolodinediones; mitoNEET; diabetes; immunological disorder; cardiovascular disorder; including hypertension; neurological disorders; ischaemia; reperfusion; ss; 2'-methoxyethyl gapmer; 2'-MOE gapmer; phosphorothioate backbone.
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                                                      WO2004053060-A2
                                                                                                                                                                                                                             Human mitoNEET-specific
                                                                                                                                                                                                                                                               09-SEP-2004
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                                                                                                                                                                                                                                                                                                                                    ADP69506 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 0 C; 1 G; 16 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2004-468836/44.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         insulin sensitising antidiabetic thiazolodinediones; mitoNEET; diabetes; immunological disorder; cardiovascular disorder; including hypertension; neurological disorders; ischaemia; reperfusion; se; 2'-methoxyethyl gapmer; 2'-MOE gapmer; phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antisense oligonucleotides encoding mitoNEET, useful for modulating ONEET expression or for treating diseases associated with mitoNEET, diabetes, immunological disorders or cardiovascular disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          1514
                                                                                      sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                        19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4; SEQ ID NO 273; 226pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense oligonucleotides encoding mitoNEET, useful for modulating mitoNEET expression or for treating diseases associated with mitoNEET, e.g. diabetes, immunological disorders or cardiovascular disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anorectic; Antidiabetic; Muscular; Gene Therapy; CpG island; IGF2 gene intron 3; muscle mass; fat deposition; teat number muscle deficiency; diabetes; PCR; primer; ss; pig.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 4; SEQ ID NO 400; 226pp; English.
     Selecting an animal for desired genotypic or potential phenotypic
                                                                                                                                                             (UYLI-)
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                                                          WPI; 2004-501307/48
                                                                                                                                                                                                                                                                                                                   10-JAN-2003; 2003EP-00075091
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                                                                                                          Andersson L,
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) MELICA HB.
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                                                                                                          Georges
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Pred. No. 4.4e+02
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RESULT 857
ADP99304/c
ID ADP993
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XX ADP993
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XW SCF/ 8
XW SCF/ 8
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(w aplastic anaemia; paroxysmal nocturnal haemoglobinuria; myelofibrosis;

(w myelosclerosis; osteopetrosis; metastatic carcinoma; acute leukaemia;

(w multiple myeloma; Hodgkin's disease; lymphoma; Gaucher's disease;

(w Miemann-Pick disease; Letterer-Siwe disease;

(w Niemann-Pick disease; Letterer-Siwe disease;

(w Niemann-Pick disease; Letterer-Siwe disease;

(w Congestive splenomegaly; Kala awar; sarcoidosis;

(w congestive splenomegaly; Kala awar; tuberculosis;

(w primary splenic pancytopaenia; miliary tuberculosis;

(w primary splenic pancytopaenia; miliary tuberculosis;

(w vitamin Bl2 deficiency; folic acid deficiency; pyridoxine deficiency;

(w vitamin Bl2 deficiency; folic acid deficiency; pyridoxine deficiency;

(w vitamin Bl2 deficiency; folic acid deficiency; pyridoxine deficiency;

(w vitiligo; neurological damage; infertility; intestinal damage;

(w vitiligo; neurological damage; infertility; intestinal damage;

(w vitiligo; neurological damage; and primer.)
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Best Local (
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                                                                               US6759215-B1
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ute leukaemia;
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Matches

Similarity

88.9%;

Score 14.8; DB 1; Pred. No. 4.4e+02; 18 T; 0 U; 0 Other;

DB 1;

Length

0

Mismatches

0

Gaps

0

Sequence 20

BP;

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1 C; 1 G;

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AAAAAAAAAAAAAAAA 1 **АААААААААААСТААА** 1537

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Coperatively linked to an expression control sequence, and isolating the Coperatively linked to an expression control sequence, and isolating the Cransformed or transfected with an expression construct comprising a CC vertebrate SCF polypeptide-encoding DNA operatively linked to a CC heterologous expression regulatory sequence, permitting the expression of the vertebrate SCF polypeptide in the host cell. Disclosed as new are rat CC and human nucleic acids encoding SCF, SCF protein fragments. The DNA CC sequences are useful for effecting the large scale synthesis of SCF by a CC variety of recombinant techniques or for generating new and useful viral CC and circular plasmid DNA vectors, new and useful viral CC and circular plasmid DNA vectors, new and useful viral CC and circular plasmid products. The DNA sequences are also useful viral CC and the related products. The DNA sequences are also useful viral CC and its related products. The DNA sequences are also useful as of protein synthesis, in genetic therapy in humans and other mammals, and CC protein synthesis, in genetic therapy in humans and other mammals, and CC paroxymmal nocturnal haemoglobinuria, myelofibrosis, multiple myeloma, CC useful for treating haematopoietic disorders, e.g., aplastic anaemia, CC paroxymmal nocturnal haemoglobinuria, myelofibrosis, multiple myeloma, CC disorders such as piebaldism and vitamin B 12 and folite acid deficiency, Diamond Blackfan anaemia, Di Guglielmo CC syndrome, congestive splenomegaly, Kala awar, sarcoidosis, primary consisting neurological damage, infertility states, intestinal damage consisting neurological damage, infertility states, intestinal damage consisting neurological damage, infertility states, intestinal damage consisted consolutes and as a consoluting from irradiation or chemotherapy, and AIDS. SCF is also useful for resulting neurological states for fighting neoplasia (cancer). The present consoluting from a miliary tubes of the present consoluting the myelocal sciency. The present consoluting the macro
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11-JUN-1990;
24-AUG-1990;
01-OCT-1990;
10-APR-1991;
25-NOV-1992;
21-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SCF) polypeptide comprising growing host cells transformed or transfected with DNA encoding a human SCF that stimulates growth of haematopoietic progenitor cells under nutrient conditions, the DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Preparing a human stem cell factor (SCF) polypeptide, useful for treating hematopoietic disorders, e.g., aplastic anemia, comprises growing host
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention relates to preparing a (vertebrate)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         transformed or transfected with DNA encoding a human SCF.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 34; 210pp; English.
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92US-00982255.
93US-00172329.
95US-00449182.
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90US-00573616.
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RESULT 858
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ADP9933
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XX SCF; a aplast
XX SCF; a separate
XX Mammal
XX Olamor
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XX V
                       The invention relates to preparing a (vertebrate) human stem cell factor (CC (SCF) polypeptide comprising growing host cells transformed or cransfected with DNA encoding a human SCF that stimulates growth of the comprising a proving host cells transformed or comprising a human SCF that stimulates growth of the comprising a human SCF that stimulates growth of comprising a proving host cells under nutrient conditions, the DNA being coperatively linked to an expression control sequence, and isolating the copyright of the comprising a construct comprising a converted comprising a converted comprising a converted comprision regulatory sequence, permitting the expression of the vertebrate SCF polypeptide in the host cell. Disclosed as new are rat combinant nucleic acids encoding SCF, SCF proteins from a number of other commission and recombinantly expressed SCF protein fragments. The DNA sequences are useful for effecting the large scale synthesis of SCF by a variety of recombinant techniques or for generating new and useful viral candidate and circular plasmid DNA vectors, new and useful transformed and crackery protein host cells, and new and useful as constituted prowth of such host cells capable of expression of CC scholad propagations. The DNA sequences are also useful as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-APR-1991;
25-NOV-1992;
21-DEC-1993;
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11-JUN-1990;
24-AUG-1990;
01-OCT-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        aplastic anaemia; paroxysmal nocturnal haemoglobinuria; myelofibrosis; myelosclerosis; osteopetrosis; metastatic carcinoma; acute leukaemia; multiple myeloma; Hodgkin's disease; lymphoma; Gaucher's disease; Niemann-Pick disease; Letterer-Siwe disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Preparing a human stem cell factor (SCF) polypeptide, useful for treating hematopoietic disorders, e.g., aplastic anemia, comprises growing host cells transformed or transfected with DNA encoding a human SCF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Stem cell factor, SCF, universal PCR primer #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-SEP-2004
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90US-00573616.
90US-00589701.
91US-00684535.
92US-00982255.
93US-00172329.
95US-00449182.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CC of protein synthesis, in genetic therapy in humans and other mammals, and Ci in developing transgenic mammalian species which may serve as eukaryotic consts for production of SCF and SCF products in quantity. The SCF is useful for treating haematopoietic disorders, e.g., aplastic anaemia, CC paroxysmal nocturnal haematopoietic disorders, e.g., aplastic anaemia, CC paroxysmal nocturnal haematopoietic disorders, in quantity. The SCF is steopetrosis, metastatic carcinoma, acute leukaemia, multiple myeloma, CC Hodgkin's disease, lymphoma, Gaucher's disease, Niemann-Pick disease, CC Hodgkin's disease, lymphoma, Gaucher's disease, Niemann-Pick disease, CC syndrome, congestive splenomegaly, Kala awar, sarcoidosis, primary CC splenic pancytopaenia, malaria, vitamin B 12 and folic acid deficiency, pyridoxine deficiency, Diamond Blackfan anaemia, and hypopigmentation CC disorders such as piebaldism and vitiligo. The SCF are also useful for treating neurological damage, infertility states, intestinal damage resulting from irradiation or chemotherapy, and AIDS. SCF is also useful for the bhancing haematopoietic recovery after acute blood loss and as a CC boost to the immune system for fighting neoplasia (cancer). The present sequence is a universal SCF PCR primer used in the isolation of SCF DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 16
                                   A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JP06303997-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                          Disclosure; Page 7; 11pp; Japanese
                                                                                                                                                                                                                                                                                    Analysis of cDNA and
                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Analysis;
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                                                                                                                                                                                                                                                                  digestion
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 restriction enzyme;
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tion enzymes.
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Sequence 21

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U; 0 Other;

Query Match

DB 1;

Length 21;

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RESULT 860
AAQ33789
ID AAQ33789
ID AAQ337
XX AAQ337
XX AAQ337
XX PCR; BE Micros
XX BOS LS
XX PCR; BE Micros
XX HOSPLIS
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                                                                                                                                                                                                                                                                                            The sequence is that of a bovine microsatellite sequence obtd. by CC screening a library of bovine MboI DNA fragments of between 250 and 500 cp with an (AC)15 and a (TC)15 oligonucleotide probe. One out of 50 cc clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (T6)n .9 microsatellites cc in the bovine genome is estimated at >100, 000. The sequence information cf for ca. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below). The sequences upstream and cdownstream of the microsatellite sequence were used to generate the required PCR primers for in vitro amplification of the corresp.

CC microsatellite (using the program OPTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of considerily important traits esp. in cattle, to allow selective breeding. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN
                                                                                                          Best Loc
Matches
                                                                                                                                                           Query Match
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                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymorphic bovine DNA markers - used in genetic identification, mapping, and selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1992-284684/34.
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02-FEB-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Georges M, Маввеу JM
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(first entry)
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                                                                                                       Score 14.8; Di
Pred. No. 4.2e-
O; Mismatches
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Pred. No. 4.2e+02;
0; Mismatches 2
                                                                                                                                    4.2e+02;
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RESULT

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AAQ75702/c
ID AAQ75702 standard; DNA; 21 BP.
XX
AC AAQ75702;
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DT 04-AUG-1995 (first entry)
XX
C Reverse transcription primer use
XX
Analysis; gene expression; rever
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Analysis; restriction enzyme; s
XX
Synthetic.
XX
Synthetic.
XX
FD 01-NOV-1994.
XX
C SYNTHET: NIPPON TELEGRAPH & TELEF
XX
PD 01-NOV-1993; 93JP-00112515.
XX
PR 16-APR-1993; 93JP-
RESULT 862
AAQ75671/C
ID AAQ756
XX AAQ756
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Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Analysis; gene expression; reverse aggregate; restriction enzyme; ss.
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    93JP-00112515.
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Pred. No. 4.2e+02;
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  Sequence
                                                                                                                                                    Disclosure; Page 7; 11pp; Japanese.
                                                                                                                                                                                             Analysis of
                                                                                                                                                                                                                                                                              16-APR-1993;
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                                                                                                                                                                              alysis of cDNA and gene expression - digestion with restriction enzymes.
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  BP;
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RESULT 865
AAQ75687/c
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AAQ75674/c
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Matches 16
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                                                                         04-AUG-1995
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aggregate; Analysis;

restriction

enzyme;

Reverse transcription primer used in cDNA analysis technique.

gene expression; reverse

transcription; primer; cDNA;

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                                                       Analysis of cDNA and gene expression - by by digestion with restriction enzymes.
                                                                                                                                                    16-APR-1993;
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Local Similarity 88.9%;
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A method for the analysis of cDNA comprises

(a) preparing an aggregate

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RESULT 867
AAQ75690/c
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                16-APR-1993;
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                                                                                                                                              aggregate;
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by digestion with restriction enzymes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                      aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ75715;
                            method can
                                                                                                                                                                                                                                  WPI; 1995-018287/03.
                                                                                                                                                                                                                                                             (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (NITE ) NIPPON TELEGRAPH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18 TAAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene expression; reverse transcription; primer; cDNA,
                            be used to analyse gene
                                                                                                                                                                                                                                                                                                                                                                                                                                       restriction
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                           93JP-00112515.
                                                                                                                                                                                                                                                                                                                        93JP-00112515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                       enzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           & TELEPHONE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.8; DB 1;
Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                            expression rapidly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORP
                                                                                                                                                                                                     by amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  by amplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
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                              and easily
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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Query Match

Sequence

21 BP; 1 A; 1 C; 2 G; 17 T; 0 U; 0 Other;

Score 14.8;

BB

۲.

Length 21;

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RESULT 871
AAQ75686/c
ID AAQ756
XX JP0630
XX AAQ756
XX AAQ756
XX WPI; 1
XAAQ756
XX AAQ756
XX AAQ76
XX 
AAQ75689/c
ID AAQ756
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Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A method for the analysis of cDNA comprises (a) preparing an aggregate (double-stranded cDNAs by using an aggregate of mRNAs and a plural type (labelled reverse transcription primers (GENESEO files AAO75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                               Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                             Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                   04-AUG-1995
                                                                                                                                                                                                                             AAQ75689
                                                                                                                                                                                                                                                                                    AAQ75689 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 1 A; 2 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 7; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Analysis of cDNA and gene expression -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1995-018287/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JP06303997-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Analysis; gene expression; reverse aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-AUG-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               digestion with restriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1519
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1519 ТАЛАЛАЛАЛАЛАСТАЛА 1536
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                                                                                                                                                                                                                                                                                                                                                                                                                           18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                        TAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93JP-00112515
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                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. 4.2e
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.8; DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e 14.0;
L. No. 4.2e+02;
--heg 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         by amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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A method for the analysis of cDNA comprises double-stranded cDNAs by using an aggregate

(a) preparing an aggregate of mRNAs and a plural type

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RESULT 873
AAQ75703/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Reverse transcription primer used in cDNA analysis technique.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1995-018287/03
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Disclosure; Page 7; 11pp; Japanese
                                          Analysis of cDNA and gene expression -
by digestion with restriction enzymes.
                                                                                                        WPI; 1995-018287/03.
                                                                                                                                                                                            16-APR-1993;
                                                                                                                                                                                                                                    16-APR-1993;
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                                                                                                                                                                                                                                                                                                                         JP06303997-A.
                                                                                                                                                                                                                                                                                                                                                                                                         aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                               Analysis; gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ75703;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 7; 11pp; Japanese.
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                                                                                                                                                  (NITE ) NIPPON
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                                                                                                                                                                                                                                       93JP-00112515.
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                                                                                                                                                                                                                                                                                                                                                                                                           enzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ВP
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No. 4.2e+02;
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); Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 21;
                                                                amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                  of mRNA followed
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RESULT 874
AAQ75705/c
ID AAQ757
XX AAQ757
XX AAQ757
XX AAQ757
XX ARPERING
DT 04-AUG
XX ARPERING
XX ARPERING
XX ARPERING
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XX JP0630
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XX O1-NOV
XX I6-APR
XX (NITE
XX WPI; 1
XX WPI; 1
XX ARPI; 1
RESULT 875
AAQ75672/c
ID AAQ756
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                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 16
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Best Local S
Matches 16
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Pred. No. 4.2e+02;
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Pred. No. 4.2e+02
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RESULT 878
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 7; 11pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Analysis of cDNA and gene expression - by digestion with restriction enzymes.
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Synthetic
                                                  aggregate;
                                                                          Analysis; gene expression; reverse transcription;
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Pred. No. 4.2e+02;
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RESULT 880
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A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798)
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                                                                                 Analysis of cDNA and gene expression - by digestion with restriction enzymes.
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; restriction enzyme; ss.
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RESULT 881
AAQ75717/c
ID AAQ7577
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TAAAAAAAAAAAAAAAA
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                                                                                                                                                                                                           BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;
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                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93JP-00112515
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                                                                                                                               1.18;
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                                                                                                         Score 14.8; DB 1;
Pred. No. 4.2e+02;
0; Mismatches 2
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Pred. No. 4.2e+02;
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                                                                                                                                                              DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
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                                                                                                            Indels
                                                                                                                                                                 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mRNA followed
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RESULT 883
AAQ75677/c
ID AAQ756
XX JP0630
XX J
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
                                                                                                                                                                                                                                            JP06303997-A
                                                                                                                                                                                                                                                                                                                                                  Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                            Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ75677
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Analysis of cDNA and gene expression - by by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-NOV-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-AUG-1995
                                                                               16-APR-1993;
                                                                                                                                                                                        01-NOV-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQ75673;
                                                                                                                                    16-APR-1993;
                              (NITE ) NIPPON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1519
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TAAAAAAAAAAAAAGTAAA 1536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 1 A; 0 C; 1 G; 19 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                           TELEGRAPH & TELEPHONE CORP
                                                                                  93JP-00112515.
                                                                                                                                    93JP-00112515
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8; D
Pred. No. 4.2e
0; Mismatches
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RESULT 884
AAQ75683/c
ID AAQ756
XX AAQ756
XX AAQ756
XX AAQ756
XX AAQ1y
E Revers
XX Analys
XX APO630
XX APO
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Best Local
                                                A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type of abbiled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                                                                                                                                                                                                                                Disclosure; Page 7; llpp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                            Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1995-018287/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Analysis; gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reverse transcription primer used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQ75683;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 7; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Analysis of cDNA and gene expression - by digestion with restriction enzymes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (NITE ) NIPPON TELEGRAPH & TELEPHONE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              restriction enzyme;
B₽;
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1 A;
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1 C; 1
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G; 18 T; 0 U; 0 Other;
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Pred. No. 4.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                             of mRNA followed
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Query Match Best Local S Matches 16

l Similarity 88.9 16; Conservative

1.1%;

Score 14.8; DB 1; Pred. No. 4.2e+02;

Length

0,

Mismatches

Indels

0

Gaps

0

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RESULT 886
AAQ75701/c
ID AAQ757
XX AAQ757
AC AAQ757
XX O4-AUG
XX Evers
XX Evers
XX Analys
KW Analys
KW aggreg
XX Analys
KW aggreg
XX Synthe
XX
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AAQ75710/c
ID AAQ757
XX AAQ757
XX AAQ757
XX AAQ757
XX ARS
XX JP0630
XX I6-APR
XX (NITE
XX (NITE
XX ARS

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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type (labelled reverse transcription primers (GENESEQ files AAO75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                Synthetic
                                                                            aggregate;
                                                                                 Analysis; gene expression; reverse transcription; aggregate; restriction enzyme; ss.
                                                                                                                                                             Reverse transcription
                                                                                                                                                                                                              04-AUG-1995
                                                                                                                                                                                                                                                                                                                     AAQ75701 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 7; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-NOV-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Analysis; gene expression; reverse transcription;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reverse transcription primer used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-AUG-1995
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16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           restriction enzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 2 A; 1 C; 1 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93JP-00112515
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                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.1%;
                                                                                                                                                             primer used
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                                                                                                                                                                                                                                                                                                                        ВP
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); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.8;
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                                                                                                                                                          cDNA analysis technique.
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. 4.2e+02;
2;
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                                                                                                            primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer; cDNA;
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Best Local S
Matches 16
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A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 1 A; 2 C; 0 G; 18 T; 0 U; 0 Other;
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                                                              Disclosure; Page 7; 11pp; Japanese
                                                                                       Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                              WPI; 1995-018287/03.
                                                                                                                                                                                 16-APR-1993;
                                                                                                                                                                                                         16-APR-1993;
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                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                           aggregate;
                                                                                                                                                                                                                                                                                                                          Analysis;
                                                                                                                                                                                                                                                                                                                                                 Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                           04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                    AAQ75709;
                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ75709
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1995-018287/03.
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                                                                                                                                                       (NITE ) NIPPON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                              gene expression; reverse
; restriction enzyme; ss.
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                                                                                                                                                                                                                                                                                                                                                                           (first
                                                                                                                                                       TELEGRAPH & TELEPHONE
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                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                          entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.8; DB 1;
Pred. No. 4.2e+02;
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                                                                                                                                                                                                                                                                                                                          transcription; primer; cDNA;
                                                                                                                                                       CORP
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                                                                                                     amplification
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                                                                                                     mRNA followed
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                             of of
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RESULT 888
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  Query Match
Best Local S
Matches 16
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                                                                                                                                                                                      The sequences represented by, AAQ90390-90401 are synthetic DNA probes containing 3' ribonucleoside termini. The sequences shown in AAQ90402-15 are synthetic DNA probes with 5' amino termini. These sequences were specific for the polymorphisms of HLA gene dQn. The sequences were used in the device of the invention. This is a self-addressable electronic device (SAED) that can be used to carry out multi-step and multiplex reactions, such as nucleic acid hybridisations. The advantages of this method are that these reactions can be carried out with complete and precise electronic control, and that the rate, specificity and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 40; 86pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New self-addressable electronic multiplex reactions such as DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1995-185870/24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Heller MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CP-1 (synthetic DNA probe with 3'ribonucleoside terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-JAN-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                  sensitivity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NANO-) NANOGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             bio:polymer synthesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1519 ТАЛАЛАЛАЛАЛАГТАЛА 1536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           \mathsf{HLA}_i dQa; 3' ribonucleoside; \mathsf{self}\text{-}\mathsf{addressable} electronic device; \mathsf{hybridisation}_i \mathsf{ss}.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18
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16; Conserv
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                                                                                                               BP; 20
                                                                                                                                                               of these reactions are greatly improved
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     Conservative
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/note= "3′ ribonucleoside terminal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                            A; 0 C; 0 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.1%;
                            1.1%;
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Score 14.8; DB 1;
Pred. No. 4.2e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          devices - used for multi-step and
hybridisation(s), clinical diagnostics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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••
                                                      Length 21;
     Indels
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AAV35395 ID AAV: XX

AAV35395 standard; DNA; 21

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RESULT 890

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ARTIOTY 889
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                                                                                                              Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                           The sequences given in ANT10742-67 are synthetic oligonucleotides which are used in the construction of the electronically self- addressable device (ED) of the invention. The ED comprises a substrate, an electrode or opt. a number of electrodes supported by the substrate, acurrent source operatively connected to the electrode and an attachment layer adjacent to the electrode which is permeable to a counterion but not permeable to a molecule capable of insulating or binding to the electrode. The attachment layer is capable of attaching a macromolecule. The ED is used for genetic typing and comprises a number of electronically addressable locations each comprising an electrode, and a binding entity, such as one of these probes, attached to each of the locations capable of detecting the presence of a genetic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Electronically self-addressable device - used for electronic control of, e.g. nucleic acid hybridisation.
                                                                                                                                                                                                                               Sequence 21 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 60; 155pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUL-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oligonucleotide
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                                                  1520 АААААААААААСТААА 1537
                                                                                                                 16;
                                                                                                                                              Similarity
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                                                                                                                 Conservative
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                                                                                                                                                                                                                               20 A; 0 C; 0 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probe, CP-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                                                                                                                     1.1%;
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"3'-ribonucleoside terminus"
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                                                                                                                 Pred. No. 4.2
0; Mismatches
                                                                                                                                           Score 14.8; DB 1
Pred. No. 4.2e+02
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                                                                                                                                                                         DB 1;
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AAZ26403
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene. The gag gene may be altered so as to produce a gag protein modified from the natural sequence by the addition, deletion or substitution of at least 1 amino acid residue. The fusion gene is inserted into a region of a vaccinia virus not essential to its propagation, to give a recombinant vaccinia virus vector which is used to transform a host cell (such as Hela, Vero, VEF, rabbit kidney RK13 or human myeloma TK-143 cella). Upon culturing the host cell produces particulate structures containing the fusion gag protein. The recombinant vaccinia virus or the fusion gag protein may be used in the production of vaccines for protecting against infection with retrovivence.
Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                      30-NOV-1999
                                                                                                                                                                                       AAZ26403;
                                                                                                                                                                                                                        AAZ26403
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 19 A; 2 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protecting against infection with retroviruses such as HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          such as HIV-1 or HIV-2, fused to a DNA fragment containing an epitope region (30-300 bases in length) of a retroviral gene other than the g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV35388-V35414 are primers used in a method which results in a retrovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 66; 84pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Recombinant vaccinia virus containing fusion HIB gag gene production in host cells of gag protein for use as vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kojima A,
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                                                                                                                   polymorphic region 592.
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                                                                                                                                                                                                                        standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                    АДАДАДАДАДА 1537
                                                                                                                                                                                                                                                                                                       AAAAAAAAAAAAAAAAA 20
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                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               region; ENV protein; vaccinia virus;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protection; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yasuda
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                                                                                                                                                                                                                                                                                                                                                                                       Score 14.8;
Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                        Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene; retrovirus;
                                                                                                                                                                                                                                                                                                                                                                                                          21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              prevention dysplastic :
                                                              Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
                                                                                                                                                                                            Human polymorphic region 415.
                                                                                                                                                                                                                                30-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 12 A; 2 C; 0 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 7; 605pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                 Homo sapiens
                                                                                                                                                                                                                                                                   AAZ26226;
                                                                                                                                                                                                                                                                                                     AAZ26226 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                   1514 TTAATTAAAAAAAAAAA 1531
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       polymorphic sites described in the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                      DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 14.8;
Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4.2e+02
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RESULT 893
AAX813
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XX IO-AUG
DT 20-AUG
DT 3' rib
XX Microe
KW Microe
KW ion-pe
KW attach
XX Synthe
XX Synthe
XX Synthe
XX W09929
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes a novel method for identifying an inhibitor CC potentially useful for treatment of cancer, where the inhibitor is active CC on a gene vital for cell growth or viability, and where the gene is CC subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is CC used for preventing the development of cancer in a patient having a CC precancerous condition, by administering to the patient a first allele CC specific inhibitor (ASI) targeted to an allele of a first sesential gene present in cells of the precancerous condition, where the normal somatic CC cells of the patient are heterozygous for the first gene, the inhibitor CC calts of the patient are heterozygous for the first gene, the inhibitor CC calts of the patient are heterozygous for the first gene, the inhibitor CC cartive on at least one but less than all allelic forms of the gene CC used in the diagnosis, prevention and treatment of LOH disorders, e.g. CC cancers, atheroscleroic plaques, premalignant metaplastic or dysplastic CC lesions, benign tumours, endometriosis, polycystic kidney disease, and CC graft versus host disease. The method can also be used to remove comover transplants. AAZZS812-ZZ6825 represent CC human polymorphic sites described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                    Microelectronic device; multi-step reaction; microscopic format; ion-permeable permeation layer; electrode; electrical control; transport; attachment; binding; DNA/RNA hybrid; probe; ss.
                                                                                                                                                                                                                                                                                 3' ribonucleoside oligonucleotide probe CP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                          20-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                  AAX81302 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21
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                                                        WO9929711-A1
                                                                                                             misc_RNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
mes 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  422 GAGTGGCGGCTGCGGCGC 439
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GAGGGGCGGCCGCCGCC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 2 A; 6 C; 12 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                               Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
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Pred. No. 4.2
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17-JUN-1999

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RESULT 894
AAV71751/c
ID AAV717
XX AAV717
XX AAV717
XX Human 15-MAR
XX HIV re
KW HIV re
KW P30 pr
XX Synthe
OS Homo s
XX HOMO s
YX II-MAR
XX II-MAR
XX (INSP
PA (INSP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          microelectronic device which is designed to actively carry out and control multi-step and multiplex molecular biological reactions in microscopic formats. A key aspect of this inventions is played by the ion permeable permeation layer which overlies the electrode. This permeation called the permit immobilization but also separates the attached oligonucleotides and hybridized target DNA sequences from the highly reactive electrochemical environment generated called the electrode surface. The microelectronic device is designed and fabricated to actively carry out and control reactions such as nucleic acid hybridizations, antibody/antigen reactions such callectronically control the transport and attachment of specific binding entities, such as nucleic acids and polypeptides, to specific microlocations. The device can subsequently control the transport and reaction of analytes or reactants at the addressed specific microlocations. The concentrate analytes and reactants, remove non-specifically bound molecules, provide stringency control for DNA hybridization reactions and improve the detection of analytes. The
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Matches
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                                                                                                                                                                                                                            HIV receptor; P30 protein; F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP;
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                                                  12-MAR-1997;
                                                                                12-MAR-1998;
                                                                                                                  17-SEP-1998
                                                                                                                                                WO9840480-A1
                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                             Human V3 loop
                                                                                                                                                                                                                                                                                                              15-MAR-1999
                                                                                                                                                                                                                                                                                                                                            AAV71751;
                                                                                                                                                                                                                                                                                                                                                                            AAV71751 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present sequence represents a probe used to exemplify the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The specification describes a self-addressable, self-assembling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 89; 179pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-385567/32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sosnowski RG, Butler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-DEC-1998;
   (INSP ) INST PASTEUR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (NANO-) NANOGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 microelectronic device designed to carry multiplex molecular biological reactions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1520 AAAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                             sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                             PHAPI;
                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                          V3 loop; human immunodeficiency virus; PHAPI; infection; therapy; PCR; primer;
                                                                                                                                                                                                                                                                             HIV receptor P30/PHAPI sense PCR primer.
                                                  97US-0040969P
                                                                                  98WO-EP001409
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.8; DB 1;
Pred. No. 4.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nerenberg MI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in m
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format.
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RESULT 895
AAZ18416
ID AAZ184
XX AAZ184
XX AZ184
XX 19-OCT
XX POlymo
XX ASTH1;
KW Lherap
XX WO9937
XX WO9937
XX 29-JUL
XX 21-JAN
XX AXYS-
PA (AXYS-
PA (AXYS-
PI Brooke
PI Miller
XX WPI; 1
XX WPI; 1
XX WAmmal
PT to dev
XX MARMIA
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression of the receptor comprises: cultivating cells transfected with a nucleotide sequence encoding P95/nucleolin, P30/PHAPI or P30/PHAPII, placed under the control of its own promoter; bringing the cells into contact with a test molecule; and quantifying expression of the P95/nucleolin, P30/PHAPI or P40/PHAPI e.g. by quantitative PCR using the primers provided (see AAV71749-54). Active molecules that have the ability to alter and/or prevent the binding of the HIV receptor to the HIV receptor can be used in pharmaceutical and diagnostic compositions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This oligonucletide is complementary to a portion of DNA sequence (see AAV71743) coding for the P30/PHAPI (see AAW84053) of the newly identified V3 loop HIV receptor. It is used as a sense primer, together with an antisense primer (see AAV71752) in a PCR amplification of P30/PHAPI reverse-transcribed mRNA. The V3 loop HIV receptor consists of an association of 3 proteins, named P95/nucleolin, P40/PHAPII and P30/PHAPI (see AAW84052-54). A method for screening molecules that modulate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated V3 loop HIV receptor - comprises P95/nucleolin, P40/PHAPII and P30/PHAPI proteins, used to develop products for the treatment and prevention of HIV infection.
                                                                 WPI; 1999-479058/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hovanessian A,
Briand J, Gui
                                Mammalian asthma related genes,
                                                                                              Brooks-Wilson AR,
Miller A, North
                                                                                                                                                                           21-JAN-1998;
                                                                                                                                                                                                            21-JAN-1998;
                                                                                                                                                                                                                                           29-JUL-1999
                                                                                                                                                                                                                                                                       WO9937809-A1
                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                  ASTH1; asthma; human; chromosome 11p; ASTH1I; ASTHIJ; genetic locus; therapeutic; immunogen; polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                  Polymorphic fragment in region 5'
                                                                                                                                                                                                                                                                                                                                                                                                                  19-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ18416 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 1 A; 10 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                            (AXYS-) AXYS PHARM INC.
                  development of asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         427 GCGGCTGCGGCGCCGGCG 444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                           98WO-US001260.
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                                                                                                               Buckler
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
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Pred. No. 4
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                                                                                                               Cardon
                                  useful for diagnosis
                                                                                                                                                                                                                                                                                                                                                                                     to ASTHLJ.
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                                                                                                                 Carey
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 21;
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                                  of a predisposition
                                                                                                                 Galvin
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RESULT 896
AAX26973/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 16
       The present primer was used to reverse transcribe RNA encoding a human mammary-specific protein, designated mammaglobin. The specification describes a protein comprising a mammaglobin antigen that is recognized by B and/or Tc cells specific for the natural, secreted and glycosylated form of mammaglobin polypeptide. This protein, or recombinant vectors that express it, are used in vaccines for treating mammaglobin are used in vaccines for treating mammaglobin expressing cancers, specifically of the breast. Such cancers can also be treated using autologous tumor lymphocytes activated ax vivo with an mammaglobin antigen, then returned to the patient. Expression of mammaglobin is elevated in 27% of stage I primary breast cancers, so it represents a marker useful for diagnosis of this disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention identifies a genetic locus ASTH1, associated with asthma, mapped to human chromosome 11p. ASTH1I and ASTHII are genes present within the locus, located close to each other on human chromosome 11p, and have similar patterns of expression, and common sequence motifs. The ASTH1 genes and fragments, encoded protein, genomic regulatory regions and anti-ASTH1 antibodies are useful in the identification of individuals predisposed to development of asthma, and for the modulation of gene activity in vivo for prophylactic and therapeutic purposes. The ASTH1 protein is useful as an immunogen to raise specific antibodies, in drug screening for compositions that mimic or modulate ASTH1 activity or expression, including altered forms of ASTH1 protein, and as a therapeutic. Sequences AAZ18366-Z18509 represent polymorphisms in the ASTH11 and ASTH1J genes
                                                                                                                                                                                                                                Example 2; Page 55; 60pp; English.
                                                                                                                                                                                                                                                                    Mammaglobin, secreted protein overexpressed
                                                                                                                                                                                                                                                                                                                                           Watson MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; mammary-specific protein; mammaglobin; antigen; vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer used to reverse transcribe mammaglobin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX26973;
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                                                                                                                                                                                                                                                                                                       WPI; 1999-244021/20
                                                                                                                                                                                                                                                                                                                                                                                                                18-SEP-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                aucorogous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mammaglobin-expressing cancer; breast cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1225 CTCTAGCTCTTAGCTTCCTC 1244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tumor lymphocyte;
                                                                                                                                                                                                                                                                                                                                         Fleming TP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diagnosis; marker; primer; ss
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Pred. No. 4.
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                                                                                                                                                                                                                                                                      in breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RNA
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                                                                                                                                                                                                                                                                       cancer
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RESULT 897
AAZ44350/c
ID AAZ443
XX AAZ443
XX AAZ443
XX AACHA
AC AARA
XX AATTAM
XX AATTAM
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XX AATTAM
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RESULT 89
AAA80318
ID AAA6
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AC AAA6
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                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 16
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     have antimisrobial, cytostatic and immunosuppressive activity. The oligonucleotides are useful for binding to and preventing or inhibiting the biological function of a protein kinase or a target molecule and for detecting the presence or absence of a target molecule in biological samples. The oligonucleotides are also useful for prophylactic and therapeutic treatment of diseases such as cancer, autoimmune diseases and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oligonucleotides inhibiting protein kinase, useful for treating such as cancer and autoimmune disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antimicrobial; cytostatic; immunosuppressive; protein kinase; prophylactic; therapy; treatment; cancer; autoimmune disease; pathogenic microorganism; primer; ss.
                           AAA80318
                                                                              AAA80318
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes novel purified aptameric oligonuclectides which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 8; Col 27-28; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-APR-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-DEC-1999
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                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      caused by pathogenic microorganisms. This sed in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
                                                                           standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
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                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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88.9%;
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Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                  Score 14.8;
                                                                                                                                                                                                                                                                                                                                             Pred.
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                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                             NO.
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                                                                                                                                                                                                                                                                                                                                                                  DB 1; Length 21;
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CC Chromosome (11p). This locus comprises the ASTHII and ASTHII genes, which CC are associated with a genetic predisposition to asthma and bronchial CC hyperreactivity. The ASTHII and ASTHII genes are oriented in opposite CC directions with the ASTHII locus, and have similar patterns of expression CC and common sequence motife. They are both expressed in trachea, lung and CC several other tissues. ASTHII and ASTHIU are novel members of the ets CC family of transcription factors, which have been implicated in the CC genes known to be important in the actiology of asthma. Both ASTHII and CASTHIU manual and activation of a variety of genes including the TCRa gene and cytokine CC ASTHIU mRNAs are alternatively spliced. Alternative splicing of asthma. Both ASTHII and CASTHIU more are all 5 to the start codon in exon b. In contrast, CC alternative splicing of ASTHII transcripts results in 3 different ASTHII concleic acids are useful as diagnostics to identify a hereditary precision of the gene in abiological specimen, and for generating genetically modified non-human animals or site specific generating genetically modified non-human animals or site specific gene inmunogens to raise specific antibodies; in drug screening for Campositions that mimic or modulate activity or expression of ASTHII candyor ASTHII and as a composition to asthma, and forms of these proteins; and as a CC antibodies are useful in the identification of individuals predisposed to cevelopment of asthma, and for modulate activity or expression of ASTHII candyor ASTHII and anti-ASTHII considered proteins, and for modulate activity in vivo for cepressent polyment of asthma, and for modulate activity in vivo for compositions or active fragments thereof, encoded proteins, and composition of asthma, and for modulation of individuals predisposed to condition of the second therapeutic purposes. The intact ASTHII or ASTHII condition of modulate or reduce bronchial hyperreactivity. Sequences AA80260-A80261 and AA80264-A80416 crepresent proteins are second.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acids other than naturally occurring chromosomes encoding ASTH1 protein, for e.g. screening compositions that modulate expression or function of ASTH1 proteins or as diagnostics for genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Galvin M, Mil
Brooks-Wilson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-JAN-1997;
01-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             splice variant;
drug screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example; Col 41-42; 131pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     predisposition to asthma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    bronchial hyperreactivity; ets family; transcription
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-NOV-2000
21
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son AR, Carey AH;
BP;
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97US-0051432P.
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9 C; 1
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G; 10 T; 0 U;
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  1 Other;
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Matches Query Match

Local

Similarity

80.0%;

1;

1.18;

Score 14.8; Pred. No. 4.

4.2e+02

DB 1;

Length 21; Indels

0

Gaps

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RESULT 899
AAP9977/C
ID AAP997
XX AAP997
AC AAF997
XX DT 12-JUN
XX Vaccin
KW Inmuno
KW Infect
XX Synthe
XX Synthe
XX 25-SEP
PR 27-SEP
RX 100A
PA (COLE-
XX UACCIN
PT USING
PT USING
PT USING
PT USING
CC The pr
CC Timmune
CC Timmune
CC Timmune
CC Thaemone
CC Th
RESULT 900
AAH42480/c
ID AAH42480 standard; DNA; 21 BP
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a staphylococcus of the present sequence can also be used to redirect a staphylococcus of the present sequence can also be used to redirect a staphylococcus of the present sequence can also be used to redirect a staphylococcus of the present sequence can also be used to redirect a staphylococcus of the sum of the present sequence can also be used to redirect a staphylococcus of the sum of the present sequence can also be used to redirect a staphylococcus of the sum of the
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27-SEP-199; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present sequence may have a phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 101; Page 56; 338pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vaccinating against tumors, infectious diseases, allergies and using immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-APR-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-JUN-2001 (first entry)
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16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                  Conservative
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99US-0156135P.
                                                                                                                                                                                                                                                                                                                                                                                                   1.1%;
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Pred. No. 4.
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RESULT 901
ABS78428/c
ID ABS784
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AC ABS784
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AC ABS784
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DT 13-DEC
XX

ABS78428;

ABS78428 standard;

DNA;

21 ВP

13-DEC-2002

(first entry)

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Matches
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Branched chain compound; nucleic acid synthesis; primer extension; reverse transcription; nucleic acid hybridization; nucleic acid amplification; ss.
                                                                                                                                                                             Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Branched compounds useful in e.g. nucleic acid synthesis reaction comprises nucleic acid moieties optionally extended by a polymera
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Schmidt W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 10; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                          The specification describes branched compounds containing nucleic acid
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                                          1520 АЛАЛАЛАЛАЛАГТАЛА 1537
21
                                                                                                              Similarity
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                                                                                         Conservative
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                                                                                                           1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "NH2-C6
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                                                                                                           Score 14.8;
Pred. No. 4.
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                                                                                         Mismatches
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                                                                                                           4.2e+02
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                                                                                                                                     DB 1;
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RESULT 902
ABL39404/c
ID ABL394
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AC ABL394
AC ABL394
AC IMMUNC
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DT 16-APR
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DT 16-APR
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeossis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scare. The present sequence is an antiangiogenic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; plaque neovascularisation; telangiectasia; haemophiliac joint; atherosclerosis;
                                                                                                                                                                           ABL39404;
                                               Antibody-induced cell lysis; cancer; immunostimulatory; CD20; anglogenesis; metastasis; cytostatic; phosphorothicate backbone; ss
                                                                                                                                                                                                             ABL39404 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Inhibiting angiogenesis in a subject, involves administering at antiangiogenic nucleic acid molecule to the subject.
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                                                                                                      Immunostimulatory nucleic acid SEQ ID NO: 840.
                                                                                                                                          16-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of the invention
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larity 88.9%;
Conservative
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                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hypertrophic scar.
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                                                                                                                                                                                                                                                                                                                                                                                                                                          A; 0 C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           precancerous lesion; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                          G; 21 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             molecule to the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           angiogenesis; solid tumour growth;
                                                                                                                                                                                                                                                                                                                                                                                       Score 14.8;
Pred. No. 4.
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RESULT 903
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                          Human; ds; cytochrome P450 A1; CYP4501A1; UGTZB4; MDR1; cytochrome P450 A2; CYP4501A2; cytochrome P450 02E; CYP45002E1; LTF; adrenergic receptor beta1; ADBR1; aryl hydrocarbon; AHR; MRP3; NR112; aryl hydrocarbon receptor nuclear translocator; ARNT; cathepsin S; CTSS cyclooxygenase 2; COX2; diazepam binding inhibitor; DBI; haematological; epoxide hydroxylase 2; EPHX2; 5-lipoxygenase activating protein; FLAP; glutathione-S-transferase 12; GST12; histamine-N-methyl transferase; HNMT; Kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT; Kallikrein 2; KLK2; nicotinamide-N-methyl transferase; NNMT; NADPH minore cytical account of the control of the control of the cytical accounts.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, breast cancer, brain and central nervous system (CNS) cancer, breast cancer, cancer, brain and cancer, stdney cancer, connective tissue cancer, oesophageal cancer, eye cancer, kidney cancer, larynx cancer, leukaemia, liver cancer, lung cancer, Hodgkin's lymphoma, non-Hodgkin's lymphoma, melanoma, myeloma, oral cavity cancer, ovarian cancer, stomach cancer, prostate cancer, rhabdomyosarcoma, skin cancer, stomach cancer, testicular cancer, and uterine cancer. The
NADPH quinone oxidoreductase 2; NQO2; sulfotransferase thermolabile; STM; UDP-glucuronosyl transferase 2B7;
                                                                                                                                                                                                    Human NADPH
                                                                                                                                                                                                                                        23-DEC-2002
                                                                                                                                                                                                                                                                                                              ABS97830 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     present sequence is an immunostimulatory oligonucleotide described exemplification of the invention % \left( 1\right) =\left\{ 1\right\} =\left\{ 1\right\} 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI;
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                                                                                                                                                                                                                                                                            ABS97830
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   developing cancer.
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                                                                                                                                                                                                quinone oxidoreductase 2 (NQO2) polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 0 A; 0 C; 0 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 309; 312pp; English.
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.1%;
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                                                                                                                                                                                                                                                                                                              В₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                    sequence #38
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Complexition relates to the sequence of an isolated nucleic acid compositing at least one base variation from that of a known human optochrome P450 A1 (CYP4501A1), Cypochrome P450 A2 (CYP4501A2), CC cytochrome P450 A2 (CYP4501A2), CYTOCHROME AND 
UGT2B7; UDP-glucuronosyl transferase; UGT2B15; urokinase receptor; uPA; multidrug resistance 1; lactotransferrin; orphan nuclear receptor; multidrug resistance associated protein 3; cancer; prostate; acetylcholine muscarinic receptor; CHWR1; CHWR2; CHWR3; CHWR4; CHWR5; altered drug metabolism; cardiovascular function; colorectal tumour; central nervous system; pulmonary; immunological; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 16; Page 130; 714pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                disorder-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Isolated nucleic acid molecules having polymorphisms in known human genes e.g. cytochrome p450 and cathepsin S useful as genetic linkage markers for locating, identifying and characterizing the genes responsible for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-698522/75.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-NOV-2000; 2000US-00724389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-NOV-2001; 2001WO-US044838.
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                                            DNA
                                            sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              traits.
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                                            the
                                            invention
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Query Match Best Local Similarity 1.1%; Score 14.8; DB 1; Length 21; Pred. No. 4.2e+02;

Sequence

21

BP;

10

A; 9

C; 1

G; 1 T; 0 U; 0 Other;

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1520 AAAAAAAAAAAAGTAAAA 1537

Conservative

<u>,</u>

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Gaps

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RESULT 904
AAD51323/c
ID AAD51323/c
ID AAD5132XX
AC AAD513
XX Lamini
XX Lamini
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XX WO2002
PD 14-NOV
XX WO2002
XX O4-NAN
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                                                                                                                                                                   performance animal. The method involves determining in sample abundance of expressed target nucleic acid; transmitting digital sample signal to remote diagnostic server; processing digital sample signal to corrented database to correlate digital signal with digital information and returning report of particular condition of animal. The method is useful for assessing a condition of a performance animal preferably human, dog or camel. The condition can be an athletic ability and a condition that enhances, hinders, impedes or does not change an expected ability of the performance animal; and also normal, pre-clinical, overt progress and/or stage of disease, undiagnosed of unclassified conditions, presence of drugs, response to exercise, response to vaccines, therapies, nutritional states and response to environmental conditions. Diseases assessed by the invention include laminitis, lameless, valiments, fractures, epistaxis, musculoskeletal damage or disorders and joint disease. The present constitutions is a primer used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Assessing condition e.g. athletic ability, stage of disease, presence of drugs, response to exercise, response to vaccines, therapies, nutritional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-MAY-2001; 2001AU-00004809
29-JUN-2001; 2001US-00896941
                                                                                                                 Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 46; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            states, of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-120558/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Brandon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-MAY-2002; 2002WO-AU000553
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Laminitis; viral disease; vaccine; bacterial disease; primer; epistaxis; gastritis; gastric ulcer; respiratory ailment; fracture; joint disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Regular oligo dT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-APR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a method for assessing a condition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 musculoskeletal damage; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GENO-) GENOMICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               545 TGTGGGTGCGTGTGCGTG 562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19 TGTGTGTGTGTGTG 2
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                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         performance animal involves analyzing nucleic acid
                                                                                                                    BP,
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                         1.1%;
                                                                                                                    0 C; 0
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Pred. No. 4.20
0; Mismatches
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                                                        Score 14.8;
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                                                                                                                       T; 0
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                                                                                                                       U; O
                               4.2e+02
                                                              DB
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                                                        Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression.
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RESULT 906
ADB37209/c
ID ADB372
XX
AC ADB372
AC ADB372
XX
DT 04-DEC
XX
DT 04-DEC
XX
Immunc
XX
GB; al
KW dB; al
KW dB; al
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OS Synthe
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ACH03246/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease.
                                                                                                                                                                                                                                                                                                                                     /209/c
ADB37209 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Treating non-allergic inflammatory diseases, such as psoriasis, allergic contact dermatitis, latex dermatitis or inflammatory by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 33; 229pp; English.
                                                                                                                 ds; allergy;
                                                                                                                                                                                                                                     04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     disease by administering an immunostimulatory nucleic acid.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAR-2001; 2001US-0279642P.
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                                                                                      hypo-responsive
                                                                                                                                                                       Immunostimulatory nucleic acid #823
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2003050268-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1520
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence represents an immunostimulatory nucleic acid
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16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAAAAAAAAAAAAAAAAAAA 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                      asthma; poly-G nucleic acid; aerosol formulation; ive subject; immunostimulatory.
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88.9%;
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Pred. No. 4.2e
O; Mismatches
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ADKO1309/c
ID ADKO13
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AC ADKO13
XX
DT 06-MA1
XX
BS; h;
KW 88; h;
KW blood
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DE Rattu
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DE RATtu
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PD 04-SE
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PD 04-SE
XX
PD 28-FI
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PR 28-FI
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PR 28-FI
PR 28
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BRAT/)
(PETE/)
(FOUR/)
                                                                         Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                            DE10208794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rattus sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rat DNA microarray
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 17; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HPI;
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                                                  and constant regions.
                                                                                                                                                    WPI; 2003-714082/68.
                                                                                                                                                                                               Boekenkamp D,
                                                                                                                                                                                                                                               (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                                                                                                               28-FEB-2002; 2002DE-01008794.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ss; hybridisation;
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FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       germ
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                                                                                                                                                                                               Dieck
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; 0 C; 0 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       capture oligonucleotide; pattern; mucosal; hair root;
cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. 4.2
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>ب</u>
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Example; Page 5; 8pp; German.

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RESULT 908
ADKO1318/c
ID ADKO13
XX ADKO13
XX ADKO13
XX AC ADKO13
XX BO ANAY
XX BO HAT DN
XX BO HAT DN
XX BO HAT DN
XX BO HAT DO
XX BO HAT DO
XX DE1020
XX BO HAT 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Comprise variable and non-variable regions. The capture oligonuclectides to least once in each nucleic acid and a 3'-variable discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids. The conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single cranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular, cardivity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible method is based on comparison of nucleic acids from cells, and can detect very small differences in pools of nucleic acid from cells, and compounds matrix miniaturisation is possible. ADK01381-ADK01344 represent conductions and the method of the invention.
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Best Local Similarity
   Sorting single-stranded nucleic acid, useful for analyzing expression
                                                                         WPI; 2003-714082/68
                                                                                                                                               Boekenkamp D,
                                                                                                                                                                                                                                                                                       28-FEB-2002; 2002DE-01008794
                                                                                                                                                                                                                                                                                                                                                                                                                                      04-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;
blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rat DNA microarray capture oligonucleotide #38.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 1 A; 0 C; 1 G; 19 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                            28-FEB-2002; 2002DE-01008794
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                                                                                                                                                                                                                        DEGUSSA BIOACTIVES
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                                                                                                                                               Dieck HT, Hoppe H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 14.8; DB 1;
Pred. No. 4.2e+02;
0; Mismatches 2
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patterns and screening active agents, uses capture agent with variable and constant regions.
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Example; Page 5; 8pp; German

Computate variable and non-variable regions. The capture oligoniclestics chave a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture comprises a sequence of 10-50, particularly 1-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, cetal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, charity of pharmaceuticals and/or nutritional compounds, e.g. food acids troot, blood, nerve or germ cells and (ii) for determining the acids troot, blood, nerve or germ cells and (ii) for determining the acids (amino, carboxylic or fatty acid) or their derivatives, ealts and mixtures. The method provides rapid, innerals, trace elements, organic cacids (amino, carboxylic or fatty acid) or their derivatives, salts and can detect very small differences in pools of nucleic acids from cells. It can detect very small differences in the nucleic acid sold. Since the complete pattern of all nucleic acid in a cell, and capture probes used in the method of the invention. reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then

Sequence 21 BP; 0 A; 1 C; 2 G; 18 T; 0 U; 0 Other;

Matches Query Match 16; Similarity Conservative 88.9%; 0 Score 14.8; DB 1; Pred. No. 4.2e+02; Mismatches DB 1; Length 21; Indels 0 0

밁 1520 AAAAAAAAAAAAGTAAAA 1537 18

ARCSULT 909
ARCSUL

ADK01323 standard; DNA; 21 ВP

06-MAY-2004 (first entry)

Rat DNA microarray capture oligonucleotide #43.

ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.

Rattus sp

DE10208794-A1

04-SEP-2003

28-FEB-2002; 2002DE-01008794

28-FEB-2002; 2002DE-01008794.

(DEGS) DEGUSSA BIOACTIVES GMBH.

Boekenkamp D, Dieck HT, Hoppe H;

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RESULT 910
ADK013
XX ADK013
XX ADK013
XX ADK013
XX B8; hy
XX B8; hy
XX B6; hy
XX B1000;
XX B1000;
XX B1000;
XX B1000;
XX B1000
XX B1000;
XX B1000;
XX B2 F2EP
XX 28-FEB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow CC binding of particular sorts of single stranded nucleic acids. The capture CC eagents are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The capture CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, regin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface by tranded nucleic acids to the specially in the form of a chip. Its surface CC is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the selectrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, thair root, blood, nerve or germ cells and (ii) for determining the actiff that the properties of patterns, especially in mucosal, CC hair root, blood, nerve or germ cells and (ii) for determining the actiff (said) or said compounds, e.g. food catiff (said) or said compounds, e.g. food catiff (said) or said (said) or their derivatives, salts and catiff (said) or said (said) or said (said) or said (said) or said (said) or catiff (said) or said (said) or said (said) or catiff (said) or said (said) or s
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Matches
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                                                                                                                                                                                                                                                                                                                          Rat DNA microarray capture oligonucleotide #64.
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  28-FEB-2002; 2002DE-01008794
                                                   28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                 Rattue ep
                                                                                                                                                                                                                                                                    ss; hybridisation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and constant regions.
                                                                                                                                                  DE10208794-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-714082/68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  comprise variable and non-variable regions. The capture oligonucleotides
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                                                                                                                                                                                                                                                    nerve;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                               tion; capture oligonucleotide; pattern; mucosal; hair root; germ cell; food additive; food supplement.
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RESULT 911
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC binding of particular sorts of single stranded mucleic acids. The capture CC agents are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-CC conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting to the surface is (quasi) covalent, supramolecular, CC physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, CC hair root, blood, nerve or germ cells and (ii) for determining the cacids (amino, carboxylic or fatty acid) or their derivatives, organic cacids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and callows imaging of the complete sapid, inexpensive and reproducible cacids in a cell, and callows imaging of the complete pattern of all nucleic acid in a cell, and companies of nucleic acid pool. Since the capture probes and in the nucleic acid pool. Since the capture probes and in the nucleic acid pool. Since the capture probes and in the nucleic acid pool. Since the capture probes in the nucleic acid pool.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local
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                                                                                                                                  ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;
blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                          Rat DNA microarray capture oligonucleotide #60.
                                                                                                                                                                                                                                                        06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example; Page 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and constant regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                   ADK01340;
                                                                                                                                                                                                                                                                                                                                               ADK01340 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   capture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1520 ААААЛААЛАЛААСТАЛАЛ 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21 AAAAAAAAAAAAAAA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                     (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.1%;
                                                                                                                                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8; Ub .,
Dred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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04-SEP-2003 DE10208794-A1

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RESULT 912
ADRO1319/c
ID ADRO13
XX ADK013
AC ADK013
XX DT 06-MAY
XX DE Rat DN
XX B9; hy
KW B100d;
XX
XX SRattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Comprise variable and or variable region, the complement of which is present at CR have a 5'-invariable anchor region, the complement of which is present at CR created a 5'-invariable anchor region of the comprises all possible combinations of up to 10 nucleotides to allow that comprises all possible combinations of up to 10 nucleotides to allow CR binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (INA) and the anchor region CR comprises a sequence of 10-50, particularly 15-25, T residues. The CR capture oligonucleotides are biotinylated and immobilized on a surface by interaction with streptavidin. The material and/or membrane, having semicant and the properties and especially in the form of a chip. Its surface by conducting properties and especially in the form of a chip. Its surface the stranded nucleic acids to the surface is (quasi) covalent, supramolecular, cphysical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, that root, blood, nerve or germ cells and (ii) for determining the carids truty of pharmaceuticals and/or nutritional compounds, e.g. food activity of pharmaceuticals and/or nutritional compounds, e.g. food carids (amino, carboxyllc or fatty acid) or their derivatives, salts and captures. The method provides rapid, inexpensive and reproducible caridows imaging of the complete pattern of all nucleic acid in a cell, and capture or permission of nucleic acid pool. Since the method is based on comparison of nucleic acid pools. Since the capture probase used in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides
      Rattus sp
                                             ss; hybridisation;
blood; nerve; germ
                                                                                                             Rat DNA microarray capture oligonucleotide #39.
                                                                                                                                                           06-MAY-2004
                                                                                                                                                                                                      ADK01319
                                                                                                                                                                                                                                               ADK01319 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example; Page 6; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Boekenkamp D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-FEB-2002; 2002DE-01008794.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                      1520 АЛЛАЛАЛАЛАД 1537
                                                                                                                                                                                                                                                                                                                                                               19
                                                                                                                                                                                                                                                                                                                                                                                                                                                  16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                             AAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dieck HT,
                                             capture oligonucleotide; pattern; mucosal; hair root;
cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hoppe H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8; DB 1;
Pred. No. 4.2e+02;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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RESULT 913 ADK01328/c

ADK01328 standard; DNA; 21

ВP

밁 S

Matches Query Match Best Local (

16;

Conservative

0;

Similarity

1.1%;

Score 14.8; DB 1; Pred. No. 4.2e+02; Mismatches

DB 1; Length 21; Indels

0

Gaps

0

SEXEXEX

ADK01328;

06-MAY-2004 (first entry)

Rat DNA microarray capture oligonucleotide #48

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CC comprise variable and non-variable regions. The capture oligonucleotides of have a 5' invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture CC agents are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The CC capture oligonucleotides are biotinylated and immobilised on a surface by CC interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-CC is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, CC physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cardistives or supplements, especially minerals compounds, e.g. food cadditives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardistives. The method provides rapid, inexpensive and reproducible method is based on comparison of nucleic acid pool. Since the capture probes used in the nucleic acid pool. Since the method of the horsential acid pools, not individual compens, matrix miniaturisation is possible. ADX01281-ADX01344 represent
Sequence 21 BP; 0 A; 2 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acids by isolation and hybridisation of nucleic acid pool reading out, where the nucleic acids are selectively bound using agents that are (a) immobilised on the surface of a solid matrix
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    patterns and screening active and constant regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-SEP-2003
                                                                                                  capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-714082/68.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (DEGS ) DEGUSSA BIOACTIVES GMBH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       describes a novel method for sorting single-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pools,
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RESULT 914
ADK01335/c
ID ADK013
XX
AC ADK013
XX
                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel method for sorting single-stranded concleic acids by isolation and hybridisation of nucleic acids pools, then compared that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at clear once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cardivity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic cardis (amino, carboxylic or fatty acid) or their derivatives, salts and cardis based on comparison of nucleic acid pools, inc acid, and cardis based on comparison of nucleic acid pools, solid in a cell, and cardis based on comparison of nucleic acid pools, not individual in the method of the invention.
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                                                                                                                                                                                                                                                                            Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                           Sequence 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Boekenkamp D,
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blood; nerve; germ cell; food additive; food supplement.
                                                                 ADK01335 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; Page 5; 8pp; German
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                                                                                                                                                                                                               1520 ААААААААААААСТАААА 1537
                                                                                                                                                                          18
                                                                                                                                                                                                                                                        l Similarity
                                                                                                                                                                                                                                                                                                                                                                                  probes used in the method of the invention.
                                                                                                                                                                          AAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                           BP; 0 A; 1 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dieck HT, Hoppe H;
                                                                                                                                                                                                                                                                          1.1%;
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                                                                                                                                                                                                                                                                          Score 14.8; DB 1; Length 21; Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                             Mismatches
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ADK01335

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creading out, where the nucleic acids by isolation and hybridisation of nucleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at cleast once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LMA) and the anchor region comprises as sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biorinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids to the surface is quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, having the cativity of pharmaceuticals and/or nutritional compounds, e.g. food cadditives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible capture probes used in the method of hucleic acid from cells. It capture probes used in the method of the invention.

CC anids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible capture probes used in the method of the invention.
                                                                                                          Query Match
Best Local
                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
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                                                                                                                                                                                                                         Sequence 21
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1520 AAAAAAAAAAAGTAAAA 1537
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                                                                                                              Similarity
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                                                                       Conservative
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                                                                                                      88.9%;
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                                                                       0,
                                                                                                          Score 14.8; DB 1;
Pred. No. 4.2e+02;
                                                                              Mismatches
                                                                                                                                              Length 21;
                                                                           Indels
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Rat DNA microarray capture oligonucleotide #22

06-MAY-2004 (first entry

ADK01302 standard; DNA; 21 BP

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                                                                                                                                                                                     CC comprise variable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture CC binding of particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The capture CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-CC conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, CC physical, stimulated by an electrical field or through a molecular sieve. CC method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the cacids (amino, carboxylic or fatty acid) or their determining the cacids (amino, carboxylic or fatty acid) or their derivatives, salts and call or detect very small differences in pools of nucleic acids from cells. It can detect very small differences in nucleic acids from cells. It can detect very small differences in nucleic acid pools, not individual composition of capture probes used in the method of the invention.
                                                                         Query Match
Best Local S
Matches 16
                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)
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                                  1520 АДАЛАЛАЛАЛАЛ 1537
                                                                                           Similarity
AAAAAAAAAAAAAAA 1
                                                                                                                                                      21 BP; 0 A; 0 C; 3 G; 18 T; 0 U; 0 Other;
                                                                           Conservative
                                                                                           88.9%;
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                                                                         0;
                                                                                           Score 14.8; DB 1
Pred. No. 4.2e+02
                                                                           Mismatches
                                                                                                                  DB 1;
                                                                                                             Length 21;
                                                                           Indels
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Matches Query Match Best Local :

Similarity

1.1%;

Conservative

<u>,</u>

Pred. No. Score 14.8;

4.2e+02

0

Gaps

0

DB 1; Length 21; Indels

Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;

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RESULT 916
RESULT 916
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ID ADKO13
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                                                       cc nucleic acids by isolation and hybridisation of nucleic acids polls, then cc reading out, where the nucleic acids are selectively bound using capture capents that are (a) immobilised on the surface of a solid matrix and (b) cc comprise variable and non-variable regions. The capture oligonucleotides cc have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture capture oligonucleotides are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, cphysical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the acids (amino, carboxylic or fatty acid) or their derivatives, salts and captures. The method provides rapid, inexpensive and reproducible captures. The method provides rapid, inexpensive and reproducible. It allows imaging of the complete pattern of all nucleic acids from cells. It captures probes used in the method of the nucleic acid pools, not individual compensive matrix miniaturisation is possible. ADK01281-ADK01344 represent to approach in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
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RESULT 917
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blood; nerve; germ cell; food additive; food supplement.
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CC comprise variable and non-variable regions. The capture oligonuclectides CC have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow comprises a sequence of 10-50, particularly 15-25, T residues. The capture CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-cc conducting properties and especially in the form of a chip. Its surface by the particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, CC physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cardity of pharmaceuticals and/or nutritional compounds, e.g. food acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardity of shaping of the complete rapid, inexpensive and reproducible method is based on comparison of nucleic acid pools, Since the method is based on comparison of nucleic acid pools, not individual components and in the mucheic acid pools. Since the method is the method of the harmonic acid pools, not individual in the method of the harmonic acid pools. Since the method is the method of the harmonic acid pools. This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) probes used in the method of the invention

Sequence 21 BP; 0 A; 0 ű N ပ္ပ 19 T; 0 U; 0 Other;

> Query Match Best Local S Matches 16 Ś 밁 WPI; 2003-714082/68 28-FEB-2002; 2002DE-01008794. 28-FEB-2002; 2002DE-01008794 04-SEP-2003 DE10208794-A1 Rattus ss; hybridisation; Rat DNA microarray 06-MAY-2004 ADK01303 standard; DNA; 21 BP Boekenkamp D, (DEGS) DEGUSSA BIOACTIVES GMBH. ADK01303 1520 ААЛАЛАЛАЛАЛАГТАЛАЛ 1537 19 ďB nerve; Similarity **AAAAAAAAAAAAAA** Conservative (first entry) germ Dieck HT, capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement. capture 1.1%; норре н; oligonucleotide N 0 Score 14.8; DB 1 Pred. No. 4.2e+02 Mismatches DB 1; #23. Length Indels 21; 0, Gaps 0;

and constant regions Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable

Example; Page 5; 8pp; German

This invention describes a novel method for sorting single-stranded cc nucleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides comprise that in each nucleic acid and a 3'-variable, discriminatory region cleast come in each nucleic acid and a 3'-variable, discriminatory region combination of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region capture oligonucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of acid, lts surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, aciditives or supplements, especially minerals, trace elements, organic cacid s(amino, carboxylic or fatty acid) or their derivatives, salts and carbox imaging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in pools of nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual

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                       have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture of capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, strimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and
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Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture bigonuclectides
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cell; food additive; food supplement.
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Pred. No. 4.2e+02;
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This invention describes a novel method for sorting single-stranded concleic acids by isolation and hybridisation of nucleic acid pools, then creading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at clear come in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture capture particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the
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germ cell; food additive; food supplement.
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Pred. No. 4.2e+02
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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25. T residues. The capture oligonucleotides are bictinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single
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RESULT 922
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Best Local :
This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid mace the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent
                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                             Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                         Example; Page 5; 8pp; German.
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cc interaction with streptavidin. The matrix is of plastic, ceramic, glass, cc metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular fillaments and binding of single crystanded nucleic acids to the surface is (quasi)covalent, supramolecular, cc physical, stimulated by an electrical field or through a molecular sieve. cc The method is used (i) for analysis of patterns, especially in mucosal, additives or supplements, egerm cells and (ii) for determining the cativity of pharmaceuticals and/or nutritional compounds, e.g. food cadditives or supplements, especially minerals, trace elements, organic cacids (amino, carboxylic or fatty acid) or their derivatives, salts and cmixtures. The method provides rapid, inexpensive and reproducible callows imaging of the complete pattern of all nucleic acid in a cell, and cc allows imaging of the complete pattern of all nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent
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                 This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region
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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, reading out, where the nucleic acids are selectively bound using car

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Example;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADK01306;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 0 A; 0 C; 2 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                  (DEGS )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1520 АДАДАДАДАДАДТАДАД 1537
                                                                                                                                                                                                                               2003-714082/68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18 AAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nerve;
                                                                                                                                                                                                                                                                                                                  DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               probes used in the method of the invention.
                                                                                                                                                                                                                                                                         Ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                           regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tion; capture oligonucleotide;
germ cell; food additive; food
                                                                                                                                                                                                                                                                         Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  capture oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21
                                                                                                                                                                                                                                                                         Hoppe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8;
Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        additive; food
                                                                                                                                                                                                                                                                         Ξ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.2e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pattern; mucosal; hair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   supplement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              root;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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RESULT 925
ADKO1325/c
ID ADKO13
XX ADKO13
XX O6-MAY
XX B8; hy
DE Rat DN
XX B8; hy
KW blood;
XX D1020
XX D21020
YX O6-FEB
XX D20020
YX OFFEB
XX OFFEB
XX OFFEB
XX OFFEB
XX Sortin
PT patter
PT patter
YX Sortin
PT patter
XX Sortin
PT patter
XX Sortin
PT patter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ঠ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC comprise variable and non-variable regions. The capture oligonuclectides have a 5' invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture CC capture oligonucleotides are biotinylated and immobilised on a surface by citeraction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal resin, gel, crystalline material and/or membrane, having semi-cc conducting properties and especially in the form of a chip. Its surface by stranded nucleic acids to the surface is (quasi)covalent, supramolecular, cphysical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the acids to the surface is (quasi)covalent, supramolecular, chair root, blood, nerve or germ cells and (ii) for determining the acids (amino, carboxylic or fatty acid) or their derivatives, salts and captures. The method provides rapid, inexpensive and reproducible or presentation of differences in pools of nucleic acid from cells. It captures robe method provides rapid, inexpensive and reproducible capture motal based on comparison of nucleic acid pools. Since the method is based on comparison of nucleic acid pools. Since the method is based on comparison of nucleic acid pools. Since the method is rate method is possible. ADX01281-ADX01344 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                             Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
                                                                                                                                                                                                                                                                                                                                                                                                          DE10208794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           blood;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ss; hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rat DNA microarray capture oligonucleotide #45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADK01325;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADK01325 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 0 A; 1 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         capture probes used in the method of the invention.
                                                                                                                                                                                           Boekenkamp D,
                                                                                                                                                                                                                                                                            28-FEB-2002; 2002DE-01008794
                                                                                                                                                                                                                                                                                                                      28-FEB-2002; 2002DE-01008794
                                                                                                                                               WPI; 2003-714082/68
                                                                                                                                                                                                                                  (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1520 AAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18 AAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nerve;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tion; capture oligonucleotide; pattern; mucosal; hair root;
germ cell; food additive; food supplement.
                                                                                                                                                                                           Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.1%;
                                                                                                                                                                                         Hoppe H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8; DB 1;
Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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Example; Page 5; 8pp; German

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable

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RESULT 926
ADK01339/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC This invention describes a novel method for sorting single-stranded concleic acids by isolation and hybridisation of nucleic acid pools, then comprise variable and non-variable regions. The capture of agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides chave a 5'-invariable anchor region, the complement of which is present at that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture capture oligonucleotides are particularly locked nucleic acids (LNA) and the anchor region comprises as sequence of 10-50, particularly 15-25, T residues. The capture of interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardives inaging of the complete pattern of all nucleic acid from cells. It can detect very small differences in the mcleic acid sfrom cells. It can detect very small differences in the mcleic acid pools, not individual capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PPXRXPXPXPXPXPXPXPXXQXXQXXQXXPXXCXI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      문
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 1 A; 1 C; 0 G; 19 T; 0 U; 0 Other;
                                                       WPI; 2003-714082/68
                                                                                                                                                                       28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                    04-SEP-2003.
                                                                                                                                                                                                                                                                                         DE10208794-A1
                                                                                                                                                                                                                                                                                                                                  Rattus
                                                                                                                                                                                                                                                                                                                                                                    ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                            Rat DNA microarray capture oligonucleotide #59.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-MAY-2004 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADK01339 standard; DNA; 21 BP
                                                                                           Boekenkamp D,
                                                                                                                                  (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                           28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1520 AAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16;
                                                                                                                                                                                                                                                                                                                                фв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity 88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                           Dieck
                                                                                           HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.8; DB 1;
Pred. No. 4.2e+02;
                                                                                             Ξ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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RESULT 927
ADK01343/c
ID ADK013
XX ADK013
XX AC ADK013
XX O6-MAY
XX B8; hy
XX B8; hy
XX B8; hy
XX B100d;
XX DE1020
XX DE1020
XX DE1020
XX 28-FEE
XX 28-FEE
XX 28-FEE
XX 1 B0eker
XX B0eker
XX B0eker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC This invention describes a novel method for sorting single-stranded CC nucleic acids by isolation and hybridisation of nucleic acid pools, then CC reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-CC is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cariotic of pharmaceuticals and/or nutritional compounds, e.g. food acids (amino, carboxylic or fatty acid) or their derivatives, salts and cariots inaging of the complete apid, inexpensive and reproducible representation of differences in pools of nucleic acid from cells. It can be method of comparison of nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix enroles used in the method of the hearth of the inventor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local (
                      Boekenkamp D,
                                                                                                     28-FEB-2002; 2002DE-01008794.
                                                                                                                                          28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                    04-SEP-2003
                                                                                                                                                                                                                         DE10208794-A1
                                                                                                                                                                                                                                                                                                      ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;
blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                Rat DNA microarray capture oligonucleotide #63.
                                                                                                                                                                                                                                                                                                                                                                                                          06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADK01343 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 0 A; 2 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       capture probes used in the method of the invention.
                                                              (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1520 AAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                        Dieck HT, Hoppe H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.8; DB 1;
Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;

supplement.

blood; nerve; germ cell; food additive; food

Rat DNA microarray capture oligonucleotide #21.

06-MAY-2004 (first entry)

ADK01301;

Rattus

28-FEB-2002; 2002DE-01008794

28-FEB-2002; 2002DE-01008794

04-SEP-2003

ADK01301/c RESULT 928

ADK01301 standard; DNA; 21

ВÞ

밁 Ś

Matches Query Match Best Local :

16;

Conservative

0;

<u>ب</u>

0,

Gaps

0

Similarity

88.9%;

Score 14.8; DB 1, Pred. No. 4.2e+02, Mismatches

DB 1;

Length 21; Indels

1520 AAAAAAAAAAAGTAAAA 1537

Sequence 21 BP; 0 A; 1 C; 0 G; 20 T; 0 U; 0 Other;

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Comprise variable and non-variable regions. The capture oligonucleotides comprise variable and non-variable regions. The capture oligonucleotides comprise variable and non-variable regions. The capture oligonucleotides comprise variable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region cc that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, cc physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucossal, cardivity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and can detect very small differences in pools of nucleic acids from cells. It can detect very small differences in the mucleic acid scid in a cell, and can detect very small differences in the mucleic acid pools. Since the complete pattern of all nucleic acid in a cell, and capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (h)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example; Page 6; 8pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-714082/68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and constant regions.
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RESULT 929
ADKO1312/c
ID ADKO13
XX ADKO13
AC ADKO13
XX DC Rat DN
XX
DE Rat DN
XX
DE Ratiod;
XX
DE Ratius
XX
DE Ratius
XX
DE Ratius
XX
DE DOd-SEF
XX
DE O4-SEF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC comprise variable and non-variable regions. The capture oligonucleotides to have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow CC binding of particular sorts of single stranded nucleic acids. The capture CC agents are particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25. T residues. The capture CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semical particularly a layer of (bio)molecular filaments and binding of single conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single conducting to the surface is (quasi) covalent, supramolecular, CC physical, stimulated by an electrical field or through a molecular sieve. CC The method is used (1) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (1) for determining the CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC mixtures. The method provides rapid, inexpensive and reproducible caids (amino, carboxylic or fatty acid) or their derivatives, salts and CC mathres in a complete pattern of all nucleic acids from cells. It can detect very small differences in pools of nucleic acids from cells. It can detect very small differences in pools, not individual complete pattern of all nucleic acids from cells, and capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                   ADK01312 standard; DNA; 21 BP
                                                                                                                                                                                                                             Rat DNA microarray capture oligonucleotide #32.
                                                                                                                                                                                                                                                                          06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 1 A; 0 C; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)
                         04-SEP-2003
                                                                    DE10208794-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                ss; hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example; Page 5; 8pp; German
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
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16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.1%;
                                                                                                                                                             capture oligonucleotide; pattern; mucosal; hair cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.8; DB 1;
Pred. No. 4.2e+02;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 21;
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RESULT 930
ADK01326/c
ID ADK013
XX
AC ADK013
XX
DT 06-WAN
XX
DE Rat DN
XX
XW 88; hy
KW 88; hy
KW 8000d;
XX
OS Rattue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC comprise variable and non-variable regions. The capture oligonucleotides (C) have a 5'-invariable anchor region, the complement of which is present at (C) least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow comprises a sequence of 10-50, particularly locked nucleic acids (LNA) and the anchor region (C) particularly locked nucleic acids (LNA) and the anchor region (C) comprises a sequence of 10-50, particularly 15-25, T residues. The capture (C) capture oligonucleotides are biotinylated and immobilised on a surface by (C) interaction with streptavidin. The matrix is of plastic, ceramic, glass, (C) metal, resin, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface (C) is particularly a layer of (bio)molecular filaments and binding of single (C) stranded nucleic acids to the surface is (quasi)covalent, supramolecular, (C) physical, stimulated by an electrical field or through a molecular sieve. (C) the method is used (i) for analysis of patterns, especially in mucosal, (C) hair root, blood, nerve or germ cells and (ii) for determining the (C) acids (amino, carboxylic or fatty acid) or their derivatives, salts and (C) acids (amino, carboxylic or fatty acid) or their derivatives, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and (C) acids (amino, carboxylic or fatty acid) or their derivatives, organic completes rapid, inexpensive and reproducible (C) acids (amino, carboxylic or fatty acid) or their derivatives, and and (C) acids (amino, carboxylic or fatty acid) or their derivatives, and (C) acids (amino, carboxylic or fatty acid) or their derivatives, and (C) acids (amino, carboxylic or fatty acid) or their derivatives, and (C) acids (amino, carboxylic or fatty acid) or their derivatives, and (C) acids (amino, carboxylic or fatty acids) or their derivatives, and (C) acids (c) acids (c) acids (c) acids (c) acids (c) a
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Best Local
                                                                                                                                       Rat DNA microarray capture oligonucleotide #46.
                                                                                                                                                                                                                                   ADK01326;
                                                                                                                                                                                                                                                                                 ADK01326 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example; Page 5; 8pp; German.
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                                                                                            ss; hybridisation;
                                                                                                                                                                                     06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Boekenkamp D,
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                                                                      nerve;
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                                                                                                                                                                                                                                                                                                                                                                                                   AAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 0 A; 0 C; 1 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                     (first entry)
                                                                   tion; capture oligonucleotide; pattern; mucosal; hair root;
germ cell; food additive; food supplement.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.1%;
88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 14.8;
Pred. No. 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          useful for analyzing expression uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 21;
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RESULT 931
ADK01305/c
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                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acids pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonuclectides
ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;
                                  Rat DNA microarray capture oligonucleotide #25.
                                                                          06-MAY-2004 (first entry)
                                                                                                                   ADK01305
                                                                                                                                                         ADK01305
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-714082/68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Boekenkamp D,
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                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   constant regions.
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                                                                                                                                                                                                                                                                                                                                                                                                                21
                                                                                                                                                         standard; DNA;
                                                                                                                                                                                                                                                        AAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                   Score 14.8; DB 1; Length 21; pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                     Mismatches
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ADK01310/ ID ADK0 XX AC ADK0 XX DT 06-M

ADK01310;

ADK01310 standard; DNA; 21

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06-MAY-2004

(first entry)

RESULT 932

밁 á

18

AAAAAAAAAAAAAAA 1

Matches Query Match Best Local S

16;

Conservative

0;

Similarity

1.1%;

Score 14.8; DB 1 Pred. No. 4.2e+02 Mismatches

DB 1;

Length 21; Indels

0

Gaps

0

1520 AAAAAAAAAAAGTAAAA 1537

Sequence 21 BP; 1 A; 1 C; 1 G; 18 T; 0 U; 0 Other;

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CC comprise variable and non-variable regions. The capture ollopoucleotides to have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture CC agents are particularly locked nucleic acids (LMA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The CC capture oligonucleotides are biotinylated and immobilised on a surface by CC interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-CC conducting properties and especially in the form of a chip. Its surface stranded nucleic acids to the surface is (quasi) covalent, supramolecular, CC physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, CC hair root, blood, nerve or germ cells and (ii) for determining the captures. The method provides rapid, inexpensive and reproducible and caids (amino, carboxylic or fatty acid) or their derivatives, salts and CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and CC andetect very small differences in pools of nucleic acid from cells. It can be the defice of the complete pattern of all nucleic acid pool. Since the method of the complete acid pools, not individual compens, matrix miniaturisation is possible. ADX01381-ADX01344 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-SEP-2003
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                                             probes used in the method
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                                             of the invention
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Rat DNA microarray capture oligonucleotide #30

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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then compared that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides chave a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region cc that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, cc metal, resin, gel, crystalline material and/or membrane, having semi-cc is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, cp physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cardial to the method provides rapid, inexpensive and reproducible acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardial based on comparison of nucleic acid gools, inceels, salts and capture nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADKO1381-ADKO1344 represent
                                                                   Matches
                                                                                  Query Match
Best Local
                                                                                                                                        Sequence 21 BP; 0 A; 0 C; 2 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-FEB-2002; 2002DE-01008794.
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blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and constant regions.
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                               1520 АААЛААЛАЛАЛАГТАЛЛА 1537
                                                                   al Similarity
16; Conserv
18
                                                                                                                                                                           probes used in the method of the invention.
AAAAAAAAAAAAAAAAA 1
                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8pp; German.
                                                                                  1.1%;
                                                                   <u>,</u>
                                                                   Score 14.8; Di
Pred. No. 4.2e
0; Mismatches
                                                                                  4.2e+02
                                                                                                      DB 1;
                                                                                                   Length 21;
                                                                     Indels
                                                                     <u>,</u>
                                                                     Gaps
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RESULT 933 ADK01336/c ID ADK01336 standard; DNA; 21 BP.

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CC reading out, where the nucleic acids are selectively bound using capture (c) reading out, where the nucleic acids are selectively bound using capture (c) comprise variable and non-variable regions. The capture oligonucleotides (c) have a 5'-invariable anchor region, the complement of which is present at (c) least once in each nucleic acid and a 3'-variable, discriminatory region (c) that comprises all possible combinations of by to 10 nucleotides to allow that comprises all possible combinations of the capture oligonucleotides are particularly locked nucleic acids (LNA) and the anchor region (c) capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, (c) metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface (c) is particularly a layer of (bio)molecular filaments and binding of single (c) stranded nucleic acids to the surface is (quasi) covalent, supramolecular, (c) hair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food (c) additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or through a molecular sieve. (c) acids (amino, carboxylic or fatty acid) or their derivatives, salts and (c) acids (amino, carboxylic or fatty acid) or their derivatives, salts and (c) acids (amino, carboxylic or fatty acid) or their derivatives, salts and (c) acids is based on comparison of nucleic acid pool. Since the method is based on comparison of nucleic acid pool. Since the complete pattern of all nucleic acid pool. Since the complete uses in the nucleic acid pool. Since the complete uses in the nucleic acid pool. Since the complete acids in the method of the invention.
                                                                                    Matches
                                                                                                                            Query Match
Best Local
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blood; nerve; germ cell; food additive; food supplement.
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                                                                                                                                                                                                                                                     Sequence 21 BP;
                                                                                                                                                                                                                                                                                                                                     capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acids by isolation and hybridisation of nucleic acid pools, then
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel method for sorting single-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example; Page 6; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-714082/68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-FEB-2002; 2002DE-01008794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADK01336;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (DEGS ) DEGUSSA BIOACTIVES GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        constant regions
1520 AAAAAAAAAAAGTAAAA 1537
                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                     0 A; 0 C; 1 G; 20 T; 0 U; 0 Other;
                                                                                                                        88.9%;
                                                                                                                                                                     1.1%;
                                                                  0;
                                                                                                                                                                Score 14.8;
                                                                                                                            Pred.
                                                                                    Mismatches
                                                                                                                            No. 4.2e+02
                                                                                                                                                                DB 1; Length 21;
                                                                                    0
                                                                                    Gaps
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RESULT 934
ADKO1342/c
ADKO1342/c
ADKO133
XX ADKO13
XX ADKO13
XX ADKO13
XX ADKO13
XX BS, h
XX CB-FEE
XX BS, h
DBCS
XX B
                                                                                                     Computies variable and non-variable regions. The capture oligonucleotides to have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow cC binding of particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25. T residues. The Capture oligonucleotides are biotinylated and immobilised on a surface by conducting properties and especially in the form of a chip. Its surface is particularly residues. The capture conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, cC physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cardivity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic cardis (amino, carboxylic or fatty acid) or their derivatives, salts and captures. The method provides rapid, inexpensive and reproducible and callows imaging of the complete spattern of all nucleic acid in a cell, and companies on function of nucleic acid pools, since the method is based on comparison of nucleic acid pools, not individual constants in the method of the hourstion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example; Page 6; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-714082/68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rat DNA microarray capture oligonucleotide #62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-MAY-2004 (first entry)
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                                                                             capture probes used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Boekenkamp D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-FEB-2002; 2002DE-01008794.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  constant regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     germ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dieck HT, Hoppe H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         capture oligonucleotide; pattern; mucosal; hair root; cell; food additive; food supplement.
                                                                                     method
                                                                                         of.
                                                                                     the invention.
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Sequence

21

BP; 0 A;

0 C; 1 G; 20 T; 0 U; 0 Other;

Local Similarity

Conservative

0

88.9%;

Score 14.8; DB 1 Pred. No. 4.2e+02 Mismatches

DB 1; Length 21; Indels

0

Gaps

0

Sequence 21 BP; 0 A; 1 C; 1 G;

19 T; 0 U; 0 Other;

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ADK01308/c
IID ADK013
XX
AC ADK013
XX
DET 06-MAY
XX
RAT DN
XX
SW blood;
XX
DE1020
XX
DE1020
XX
DE1020
XX
O4-SEP
PM 04-SEP
PM 28-FEB
XX
CDEGS
PM (DEGS
XX
MPI; 2
XX
Boeken
XX
WPI; 2
XX
CC This i
CC 
                                                             This invention describes a novel method for sorting single-stranded concleic acids by isolation and hybridisation of nucleic acid pools, then complete that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow comprises a sequence of 10-50, particularly 15-25, Tresidues. The capture CC agents are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, Tresidues. The capture crapture oligonucleotides are bickinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cativity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic cards (amino, carboxylic or fatty acid) or their derivatives, salts and concentratives. The method provides rapid, inxpensive and reproducible can detect very small differences in pools of nucleic acid from cells. It genes, matrix miniaturisation is possible. ADK01281-ADK01344 represent trace persons used in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADK01308;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example; Page 5; 8pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and constant regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-714082/68.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rat DNA microarray capture oligonucleotide #28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-MAY-2004 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1520 AAAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 АДАДАДАДАДАДАДАДА
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tion; capture oligonucleotide; pattern; mucosal; hair root;
germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21
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Score 14.8; DB 1;

Length 21;

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RESULT 936
ADKO1311/c
ID ADKO13
XX ADKO13
XX ABA DAX
DT 06-MAY
XX 88; hy
XX 88; hy
XX Blood;
OS Rattus
XX DE1020
XX DE1020
XX O4-SEP
XX O4-SEP
XX (DEGS
PN DE1020
XX PAT
EN BOCKET
XX SORTH
PT Patter
PT Patter
PT Patter
CC Compri
CC Compri
CC That (CC Compri
CC Compr

CC nucleic acids by isolation and hybridisation of nucleic acids aby isolation and hybridisation of nucleic acid pools, then CC agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonucleotides CC least once in each nucleic acid and a 3 -variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture CC apture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio) molecular filaments and binding of single stranded nucleic acids. The surface is particularly a layer of (bio) molecular filaments and binding of single combinated by an electrical field or through a molecular, by stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in nucleic acids for the method provides rapid, inexpensive and reproducible and complements, especially minerals, trace elements, organic cardes (amino, carboxylic or fatty acid) or their derivatives, salts and cardinares. The method provides rapid, inexpensive and reproducible is allows imaging of the complete pattern of all nucleic acid single it the mothod is based on comparison of nucleic acid pools, not individual can be complete pattern of all nucleic acid pools. Since the cardinares matrix miniaturisation is possible. ADK01281-ADK01344 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ঠ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example; Page 5; 8pp; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;
blood; nerve; germ cell; food additive; food supplement.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel method for sorting single-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-714082/68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-FEB-2002; 2002DE-01008794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DE10208794-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rat DNA microarray capture oligonucleotide #31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADK01311;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-FEB-2002; 2002DE-01008794
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CC have a 5'-invariable anchor region, the complement of which is present at CC that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorte of single stranded nucleic acids. The capture CC binding of particularly locked nucleic acids (LNA) and the anchor region CC comprises a sequence of 10-50, particularly 15-25, T residues. The CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semi-CC conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single crystalline material and/or membrane, having semi-CC physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, hair root, blood, nerve or germ cells and (ii) for determining the CC acids (amino, carboxylic or fatty acid) or their derivatives, salts and compounds, e.g. food cacids (amino, carboxylic or fatty acid) or their derivatives, salts and compounds. The method provides rapid, inexpensive and reproducible constitution of differences in pools of nucleic acids from cells. It
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blood; nerve; germ cell; food additive; food supplement.
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Pred. No. 4.2e+02;
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                              CC This invention describes a novel method for sorting single-stranded CC nucleic acids by isolation and hybridisation of nucleic acid pools, then CC reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture comprises are particularly locked nucleic acids (LNA) and the anchor region CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, certail, resin, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly 1 alayer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, charity of sharmacation, nerve or germ cells and (ii) for determining the certails of particular the control of the contro
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Pred. No. 4.2e+02;
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                    This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resid, gel, crystalline material and/or membrane, having semi-conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single
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germ cell; food additive; food supplement.
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The invention relates to a method of enriching a low abundance polynucleotide in a sample of polynucleotides comprising a low abundance and a high abundance polynucleotide. The method comprises exposing the sample to an enzymatically non-extendable nucleobase oligomer having a nucleobase sequence complementary to a sequence within the high abundance polynucleotide under conditions so that base pairing occurs, and subjecting the sample to conditions for polymerase extension. Preferably, the enzymatically non-extendable nucleobase oligomer does not have a ribose-containing oligomeric structure. It is a peptide nucleoc acid (PNA) oligomer or is a modified nucleotide oligomer or internucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADM96310
                                                                                                                                                                                                                                            Enriching low abundance polynucleotides in a sample, useful for gene expression analysis, comprises exposing the sample to an enzymatically non-extendable nucleobase oligomer to block polymerase activity on high
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Coligomers. The modified nucleotide oligomer or internucleotide analogue coligomers is selected from locked nucleic acids (LNA), N3'-P5' Coligomer is selected from locked nucleic acids (LNA), N3'-P5' CC oligomers (MB-linked oligomers, minor groove binder-linked-coligomers, C1-C4 alkylphosphonate oligomers, phosphoramidates, beta-cc oligomers, C1-C4 alkylphosphonate oligomers, phosphoramidates, beta-cc oligomers, C1-C4 alkylphosphonate oligomers, phosphoramidates, beta-cc oligomers is chimeric. The care enzymatically non-extendable nucleobase oligomer is chimeric. The sample comprises more than one high abundance polynucleotide. The sample comprises more than one high abundance comprises transcription to yield a first strand cDNA. The method further comprises more than one high abundance comprises second strand cDNA synthesis. The sample is exposed to the nucleobase oligomer during the first and/or second strand cDNA synthesis. The sample is exposed to the comprises an amplification step, which is by creverse transcription (PCR) or by in vitro transcription. The RNA is cmand of cRNA or cotal cellular RNA. Alternatively, the sample comprises CC DNA, and polymerase extension is by DNA-dependent DNA polymerase in a comprises labelling the amplification. The cc methods are useful in selective enrichment of low abundance polynucleotides in a sample. The pool of enriched polynucleotides may be used in analysing gene expression and in creating cDNA libraries. The cc presents sequence represents a reverse transcriptiae (RT)-PCR primer which cannsporting, mitochondrial ATP synthase, subunit B, isoform 1 (ATP5F1) cc gene.
    RESULT 941
ABD25907
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                   Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                analogue oligomer. The modified nucleotide oligomer is selected from 2'-modified and 3'-modified nucleotide oligomers. The 2'-modified and 3'-modified nucleotide oligomers are selected from 2'-O-alkyl modified nucleotide oligomers and 3'-alkyl modified nucleotide oligomers. The 2'-O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABD25907;
                                        24-APR-2001; 2001US-0286036P
                                                                                  23-APR-2002; 2002WO-US013143
                                                                                                                                                                       WO200285309-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                      AI654215-derived oligonucleotide SEQ ID 4919.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABD25907 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -alkyl modified nucleotide oligomers are 2'-O-methyl nucleotide
(EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21 AAAAAAAAAAAAAAAA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAAAAAAAAAAGTAAAA 1537
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; 0 C; 0 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.8;
Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      8; DB 1;
4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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RESULT 942
ADJ88057/c
ID ADJ880
XX ADJ880
XX ADJ880
XX ADJ880
XX ADJ880
XX BA CALL
XX BA CALL
XX Select
KW Select
KW Select
KW Select
KW Select
XX Uniden
XX US2004
XX US2004
XX US2004
XX ADJ88057/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC comprising oligonucleotides, effective for alleviating controlled the sensitivity, levels of adenosine (A) or (A) receptors, construction, respiratory tract inflammation, allergies and concentration of hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating controlled the target polypeptide associated with lung airway or lung controlled the sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition composition composition of availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasconstriction, respiratory chairs, impeded respiration, respiratory condition, allergies, asthma, impeded respiration, respiratory condition, chronic obstructive pulmonary disease, pulmonary conditions to condition such as pulmonary disease, pulmonary conditions to condition the condition of the condition are associated conditions are associated conditions. The condition condition is condition to the condition are associated conditions are associated conditions. The condition is condition are associated conditions are associated conditions are associated conditions. The condition is condition are associated conditions are associated conditions are associated conditions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, t prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                       Selective enrichment;
                                                                                                                                                                                               RT primer used in the synthesis
                                                                                                                                                                                                                                             06-MAY-2004
                                                                                                                                                                                                                                                                                                                                  ADJ88057
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 15; SEQ ID NO 4919; 763pp; English.
  22-JAN-2004
                                             US2004014105-A1
                                                                                        Unidentified
                                                                                                                                                                                                                                                                                        ADJ88057;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes a novel composition (a) a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1520 АААААААААААСТАААА 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21
                                                                                                                                                                                                                                                                                                                                  standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Li Y, San
Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                             AAAAAAAAAAAAAAAA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sandrasagra A,
, Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; 0 C; 0 G; 0 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                     gene
                                                                                                                                                                                                                                                                                                                                  2
                                                                                                                                                       expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.8; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Katz
S;
                                                                                                                                                                                               of an artificial gene transcript
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.2e+02
                                                                                                                                                       RT; reverse transcriptase; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pabalan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 21;
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RESULT 943
ADM07216/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to methods for the selective enrice to abundance polymucleotides. The invention is useful for analy expression in a sample and creating cDNA libraries. The present is reverse transcriptase (RT) primer used in the synthesis of a artificial gene transcript.
                                                          Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Enriching low abundance polynucleotide relative to a high abundance polynucleotide in a sample, for analyzing gene expression and creating cDNA libraries, comprises blocking polymerase activity on high abundance
                                                                                                                                                                                                                                         Double-stranded cDNA synthesis; cDNA first strand synthesis; cDNA second strand synthesis; RNA template; RNA amplification;
                                       WPI; 2004-340131/31.
                                                                                        (SCHR/)
                                                                                                                                                  23-OCT-2002; 2002US-00278760
                                                                                                                                                                                         US2004081962-A1
                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                          15-JUL-2004
                                                                                                                                                                                                                                                                                                                                 ADM07216 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 0 A; 0 C; 0 G; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-121562/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Schroeder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-MAY-2002; 2002US-00144179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-MAY-2003; 2003US-00435489
                                                                                                           (CHEN/)
                                                                                                                             23-OCT-2002; 2002US-00278760
                                                                                                                                                                      29-APR-2004
                                                                                                                                                                                                                                                                                                              ADM07216;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; SEQ ID NO 41; 62pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 polymucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CHEN/)
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                                                          ú
                                                                                                                                                                                                                                                                                                                                                                                   21
                                                                            ) CHEN C.
) SCHROEDER I
) BRANDIS J.
) SCHROTH G.
                                                                                                                                                                                                                                                                                                                                                                                                                          16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ) SCHROEDER B G
) CHEN C.
) SCHROTH G P.
                                                                                                                                                                                                                                                                        primer used
                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                          Schroeder B,
                                                                                                                                                                                                                                                                                                                                                                                   AAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BG,
                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                                                 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chen
                                                                                                                                                                                                                                                                                                                                  DNA;
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                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                   1.1%;
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                                                           Brandis
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                                                                                                                                                                                                                                                                                                                                   ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                        first
                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.8; DB 1.
Pred. No. 4.2e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T; 0
                                                                                                                                                                                                                                                                        strand synthesis.
                                                           Schroth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ς,
                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                           ဂ္
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the selective enrichment of is useful for analysing gene
                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence
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0

Synthesizing double-stranded cDNA, by synth template, removing the template and synthes using the cDNA as template in the presence

by synthesizing a cDNA strand from R d synthesizing double-stranded cDNAs presence of processive DNA polymerase

random

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RESULT 944
ADP09287/c
ID ADP092
XX ADP092
XX ADP092
XX Extend
XX Extend
XX Becret
XW Single
XX Homo 8
XX Homo 8
XX 10-JUN
XX 25-NOV
YR 25-NOV
YR 25-NOV
PR 24-JUL
XX Roth R
XX Roth R
XX Identi
PT or abs
PT diagno
XX Identi
PT or abs
PX Exampl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC primer complementary to template, removing the template, Synthesising CC double-stranded cDNAs in a second reaction mixture comprising processive CDNA polymerase, DNA ligase, first cDNA strand as template and random CC primers having a mixture of oligonuclectides having random DNA sequences. CC Also disclosed is a method for amplifying a population of RNA molecules CC to produce a pool of double-stranded cDNA molecules, and a kit for CC synthesising quantitative information about the genetic profile of nucleic acid in original RNA sample. The method of the invention is CC useful in differential gene expression assays for the analysis of CC useful in differential gene expression assays for the analysis of CC diseased and normal tissue and for large-scale correlation studies on CC sequences, mutations, variants or polymorphisms among samples. The method CC is efficient in synthesising improved cDNA molecules and effective in CC generating useful quantities of an amplified cRNA product that comprises CC a population of cRNA molecules in substantially the same relative molar cratio as the RNA or mRNA starting material. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            breast cancer; cytostatic; gene therapy; human; chromogranin B; secretogranin 1; SCG1; chromosome 20pter-p12; ss; PCR; primer; ssingle nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to a method for synthesising double-
stranded cDNA, by synthesising first cDNA strands in a first reaction
mixture comprising reverse transcriptase, RNA template, and first strand
                      Example 5; Page 103; 286pp; English
                                                       Identifying a subject at risk of breast cancer by detecting the presence or absence of one or more nucleotide polymorphic variations, useful for diagnosing, preventing and/or treating breast cancer.
                                                                                                                                                                                                                                                         25-NOV-2002;
24-JUL-2003;
                                                                                                                                                                                                                                                                                                                   25-NOV-2003; 2003WO-US037966
                                                                                                                                                                                                                                                                                                                                                                                                  WO2004047767-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Extend primer 82 used to genotype human chromogranin B polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADP09287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                       WPI; 2004-441082/41.
                                                                                                                                                                                                                                                                                                                                                              10-JUN-2004.
                                                                                                                                                                                                                    (SEQU-) SEQUENOM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAAAAAAAAAAGTAAAA 1537
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                                                                                                                                                                           Nelson MR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                         2002US-0429136P.
2003US-0490234P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                  Kammerer SM, Reneland
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RESULT 945
ABL60935
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Best Local S
Matches 16
                                 Matches
                                                Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel method for identifying a subject at risk of breast cancer which comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The method of the invention has cytostatic applications and may be useful for identifying a risk of breast cancer, as well as therapeutic and prophylactic treatments that specifically target breast cancer, such as gene three current sequence is that of an Extend primer of the invention which was used to genotype single nucleotide polymorphisms within human chromogranin B (CHGB;secretogranin 1;SCG1) DNA which is located at chromosomal position 20pter-p12.
                                                                                                                      The invention relates to a novel human nucleotide reducing enzyme 59.62 polypeptide and encoding polypucleotide. The polypucleotide, polypeptide and its antagonist are useful for treating e.g. embryo development teratogenesis, blood system disease, and growth development disturbance disease. The present sequence represents the human nucleotide reducing enzyme 59.62 cDNA isolating RT-PCR primer
                                                                                                                                                                                                                                                                     Human nucleotide reducing enzyme 59.62 polypeptide and its encoding polynucleotide, for treating e.g. embryo development teratogenesis.
                                                                                             Sequence 24
                                                                                                                                                                                                                                                                                                                         WPI; 2002-305607/35
                                                                                                                                                                                                                                                                                                                                                        Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                      07-JUL-2000; 2000CN-00117037
                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-JUL-2000; 2000CN-00117037
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human nucleotide reducing enzyme 59.62 cDNA isolating
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                                                                                                                                                                                                                                            Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  system disease; human; RT-PCR; primer; ss
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                                                 Similarity
   TGTTTTGTTTTTAATCA 1267
                                                                                                                                                                                                                                          Page 17 (disclosure); 34pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   reducing enzyme 59.62; embryo development;
                                 Conservative
                                                                                              BP; 6 A; 2 C; 2 G; 14 T; 0 U; 0 Other;
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88.9%;
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                                                Score 14.8; DB 1;
Pred. No. 3.6e+02;
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Pred. No. 4.2e+02;
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RESULT 947
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ID ADM164
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ARTS9640/c
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Matches 15
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                                             ADM16445;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Measuring telomerase activity, useful for tumour diagnosis and compound screening - by extending substrate primer, followed by amplification and immobilising product for detection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Telomerase; substrate; primer; detection; 5'-region; retrovirus; long terminal repeat 2; LTR-2; diagnosis; tumour; screening; effector compound; PCR; amplification; Oligo-dT-Primer; ss.
    17-JUN-2004
                                                                                     ADM16445
                                                                                                                                                                                                                                                                                                                                                             Sequence 19
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les 15; Conserv
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                                                                                       standard;
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Pred. No. 5.1e+02;
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AAQ75722;

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Matches
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                                                                                                                                                                                                                                                      The invention relates to isolated RNAs comprising an intron RNA that is released in a cell, thus modulating the function of a target gene. Also disclosed is a DNA template for the isolated RNA, an expression vector comprising the DNA, and a composition comprising one or more agents that induce RNA-mediated modulation of the functions of two or more target genes in a cell, such as a mammalian cell. The isolated RNAs and compositions are useful for modulating the function of a target gene in a cell, e.g. to inhibit a cancer-related gene, potential viral gene, and microbe-related gene, and thus useful for treating and preventing diseases such as cancer and viral/microbial infections. The current sequence represents a potential poly-pyrimidine tract of the artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated RNAs comprising an intron RNA that is released in a cell, thus modulating the function of a target gene, useful for treating and preventing diseases such as cancer and viral/microbial infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cytostatic; antimicrobial; virucide; gene therapy; RNA intron; cancer; viral; microbial; infection; poly-pyrimidine tract; ds.
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12-OCT-2002; 2002US-0418405P.
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                                                                                                                                                                                          Sequence 19 BP; 0 A; 3 C; 0 G; 0 T; 13 U;
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                                            1520 ААЛААЛАЛАЛАДСТАЛА 1536
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RESULT 949
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                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                        Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
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                              WPI; 1995-018287/03
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                                                                                 (NITE ) NIPPON TELEGRAPH & TELEPHONE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     restriction enzyme; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry
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                                                                                                                                   93JP-00112515.
                                                                                                                                                                                         93JP-00112515
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.6; DB 1; Length 21;
Pred. No. 4.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in cDNA analysis technique.
                                                                                    CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cDNA analysis technique.
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RESULT 950
AAQ75734
ID AAQ757
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Best Local
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                                                                                A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GRNESEQ files AQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                           Sequence 21
                                                                                                                                                                                                                                                                                             WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                 16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                                                                                                                    Disclosure; Page 8; llpp; Japanese.
                                                                                                                                                                                                                                                  by digestion with restriction
                                                                                                                                                                                                                                                              Analysis of cDNA and gene expression -
                                                                                                                                                                                                                                                                                                                        (NITE ) NIPPON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ75734 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21
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1 Similarity
17; Conserv
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17; Conservative
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   Conservative
                                                           BP; 2 A; 2 C; 0 G; 17 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                     93JP-00112515.
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             1.0%;
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Pred. No. 4.
               Score 14.6; DB 1;
Pred. No. 4.6e+02;
                                                                                                                                                                                                                                                  enzymes
   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                         CORP
                                                                                                                                                                                                                                                              λq
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 21;
                                                                                                                                                                                                                                                                amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amplification
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                            Length
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   Indels
                              21;
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 Gaps
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1246 TCTTTGTTTTGTTTTTAATC 1266

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RESULT 951
AAQ75720
ID AAQ757
XX AAAUG
XX AAAUG
XX JP063(
YA JP063
ADKO1285
ID ADKO12
XX ADKO12
XC ADKO12
XX ACCOMAN
ACCOMAN
ACCOMAN
XX B8; hy
KW B100d;
XX
XX B100d;
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XX B100d;
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XX DE1020
XX
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A method for the analysis of cDNA comprises (a) preparing an aggregate (a) double-stranded cDNAs by using an aggregate of mRNAs and a plural type (labelled reverse transcription primers (GENESEQ files AAO75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                           DE10208794-A1
                                                                                            Rattus sp.
                                                                                                                                                                                      se; hybridisation;
                                                                                                                                                                                                                                          Rat DNA microarray capture oligonucleotide #5.
                                                                                                                                                                                                                                                                                                           06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                     ADK01285
                                                                                                                                                                                                                                                                                                                                                                                                                                ADK01285 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 8; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Analysis; gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-AUG-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1302 TCTATTTTTTTTATTTTCAGA 1322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 μ
                                                                                                                                                          nerve;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 3 A; 0 C; 1 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                       germ
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                                                                                                                                                       capture oligonucleotide; pattern; mucosal; hair root;
cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.0%;
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Pred. No. 4.6e+02;
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This invention describes a novel method for sorting single-stranded conductive acids by isolation and hybridisation of nucleic acid pools, then comprise variable then nucleic acids are selectively bound using capture comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at certain once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow comprises are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids. The conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, computed by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, having semical sused (i) for analysis of patterns, especially in mucosal, cards (amino, carboxylic or fatty acid) or their derivatives, salts and cards (amino, carboxylic or fatty acid) or their derivatives, salts and cards (amino, carboxylic or fatty acid) or their derivatives, salts and can detect very small differences in pools of nucleic acid from cells. It can detect very small differences in the mucleic acid from cells. It capture probes used in the method of the invention.
                                                                                                                                                                                                                                          RESULT 953
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                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local :
                  ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.
                                                                                                                       06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-SEP-2003
                                                                              Rat DNA microarray capture oligonucleotide #15.
                                                                                                                                                            ADK01295;
                                                                                                                                                                                                   ADK01295 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                            1302 TCTATTTTTTTTATTTTCAGA 1322
                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                       (first
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                                                                                                                                                                                                   DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.6;
Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                    4.6e+02
                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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DE10208794-A1

Rattus

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RESULT 954
AAQ75762
ID AAQ757
XX
AC AAQ757
XX
DT 04-AUG
XX
DE Revers
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                                                                                                                                                                                                                                                                                                                                                              This invention describes a novel method for sorting single-stranded CC nucleic acids by isolation and hybridisation of nucleic acid pools, then CC reading out, where the nucleic acids are selectively bound using capture CC agents that are (a) immobilised on the surface of a solid matrix and (b) CC comprise variable and non-variable regions. The capture oligonucleotides CC have a 5'-invariable anchor region, the complement of which is present at CC least once in each nucleic acid and a 3'-variable, discriminatory region CC that comprises all possible combinations of up to 10 nucleotides to allow CC binding of particular sorts of single stranded nucleic acids. The capture CC agents are particularly locked nucleic acids (LMA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The CC capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, CC metal, resin, gel, crystalline material and/or membrane, having semiconducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is quasil covalent, supramolecular, chair root, blood, nerve or germ cells and (ii) for determining the activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and compounds imaging of the complete pattern of all nucleic acid from cells. It can detect very small differences in the mucleic acid good. Since the method is based on comparison of nucleic acid pools, not individual capture probes used in the method of the invention.
                                                                                                                                                                                                          밁
                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
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                                     04-AUG-1995
                                                                                                              AAQ75762 standard;
                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                   (first entry)
                                                                                                                                                                                                                                                                                     Conservative
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Reverse transcription primer used in cDNA analysis technique

Analysis of cDNA and gene expression – by amplification of mRNA followed by digestion with restriction enzymes.

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RESULT 955
AAQ75634
ID AAQ756
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Best Local S
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                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                               Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
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WPI; 1995-018287/03.
                                                                                          16-APR-1993;
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                                           (NITE ) NIPPON
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17; Conserv
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                                              TELEGRAPH & TELEPHONE CORP
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Pred. No. 4.6e+02
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RESULT 956
AAQ75682
ID AAQ756
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Best Local Similarity
Matches 17; Conserv
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                                                                                                                                                                                                                                                                                                      A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AQ075547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            digestion with restriction enzymes.
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17; Conserv
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                                                                                                                                                                                                                        BP; 1 A; 1 C; 0 G; 19 T; 0 U; 0 Other;
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Pred. No. 4.6e+02;
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Pred. No. 4.6e+02;
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                                                                                                                                                             DB 1;
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16-APR-1993;
                           01-NOV-1994
                                                                             Synthetic.
                                                                                                     aggregate;
                                                                                                                  Analysis; gene expression; reverse transcription; primer; cDNA
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                                                                                                                                                                                                                         AAQ75764 standard;
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                                                                                                                                          Reverse transcription primer used in cDNA analysis technique.
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         A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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          Reverse transcription primer
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  Disclosure; Page 7; 11pp; Japanese
                            Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                            16-APR-1993;
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Pred. No. 4.6e+02;
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                        The present invention describes a method for the comparative detection of the amount of an RNA. The method comprises; (a) cDNAs obtained by compared by using a same restriction enzyme; (b) each different adaptor and a common adaptor are added to each of the cDNA fragments derived from the same or different tissues by the step (a); (c) the resultant adaptor added cDNAs are mixed together; (d) an adaptor primer having the common sequence to said different adaptor and a gene-specific adaptor are used to amplify said adaptor-added cDNAs containing no region derived from compositive acid of the mRNA before the addition of the adaptor among the adaptor-added cDNAs prepared by the step (b); (e) the ratios of the CDNA amounts are measured between the tissues; (f) the RNA is detected from the measured result; (g) each different adaptor and a common adaptor comitive transfers; (i) the adaptor-added genomic DNAs are amplified by using an adaptor primer having the common sequence to the different adaptor and a sequence-specific adaptor; and (j) the ratios of the amplified by using of the genomic DNAs are measured between the tissues; the ratios of the amplified mounts of the genomic DNAs are amplified by using the common sequence to the different adaptor and a sequence-specific adaptor; and (j) the ratios of the amplified mounts of the genomic DNAs are measured between the individuals. The method is used for the detection of the amounts of RNA and DNA. The present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 9; 9pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Comparative detection of the amounts of RNA and DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-MAY-2000; 2000JP-00160324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-MAY-2000; 2000JP-00160324.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-DEC-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PolyA adaptor oligonucleotide SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           comparative detection; adaptor; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
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Pred. No. 4.
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RESULT 964
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XX Human;
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention provides the protein and coding sequences of human phosphatidylinositol-3 (PTDINS-3) kinase 35. The sequences can be used in the treatment of cancer, haemopathy, HIV infection, development disorders, immunological diseases and inflammation. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New human phosphatidylinositol-3 (PTDINS3) kinase 35 for diagnosing and treating malignant tumor, hemopathy, human immunodeficiency virus infection, immunological diseases and various inflammations.
                         24-SEP-2002
                                                                             ABK86169;
                                                                                                                            ABK86169 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; Page 12; 34pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2002-025836/03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; phosphatidylinositol-3 kinase 35; PTDINS-3 kinase 35; cancer; haemopathy; development disorder; HIV infection; immunological disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human phosphatidylinositol-3 kinase 35 cDNA PCR primer #2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      is a PCR primer for the coding sequence of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-MAR-2001; 2001WO-CN000328
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                                                                                                                                                                                                                                                                                                                                                   \text{Natch 1.0%; Local Similarity 81.0%; Local Similarity 81.0%; test 17; Conservative
                                                                                                                                                                                                                                                                                                    1302
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 3 A;
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                         (first entry)
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                                                                                                                            DNA;
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                                                                                                                                                                                                                                                     CTTTTTTTAAGA 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 C; 1 G; 20 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                             Score 14.6;
Pred. No. 4e
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Pred. No. 4
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                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 24;
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Matches Query Match Best Local

al Similarity 81.0

81.0%;

Score 14.6; DB 1; Mismatches

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1246 TCTTTGTTTTGTTTTTTTAATC 1266

21

TTTTTTTTTTTTTTTAAAC

Oligo dT primer #1 used in

method

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study gene expression

24-SEP-2002 ABK86168 ABK86168

(first entry)

standard; DNA; 24 BP

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CC of gene expression in one or more samples, comprising an identifier having a specific oligo-dT primer sequence, where the identifier comprises a CC detectable marker at its 5' end. The system is useful for identifying any detectable marker at its 5' end. The system is useful for identifying any CC or all genes expressed in a given in vivo or in vitro RNA sample, as well can sthe relative differences in mRNA between 2 or more samples, where CC desired, for supporting discovery of new genes, and for identifying mRNAs that are expressed in a different levels between 2 or more samples. The new CC system or method addresses limitations of prior methods by comprising CC compositions and systems that incorporate new strategies where molecular or biochemical assay compositions and systems are linked to DNA or RNA cc sequence databases for optimal resource efficiency in assaying gene CC expression. The system has the following advantages over existing CC methods: (a) prior sequence information or clone library construction is not needed to enable the assay; (b) provides immediate sequence information in addition to information concerning changes or differences in mRNA level, to determine mRNA expression level and mRNA identification in one assay; (c) generates cDNA fragments from all mRNAs present in the cc sample for subsequent investigation by common molecular biology compositions lacking significant genomic sequence in formation. The present consequence represents an oligo dT primer used in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying and characterizing gene expression in samples, identifying mRNAs expressed at different levels, comprises identimer having a oligo-dT primer of a specific sequence a detectable marker at its 5' end.
Sequence 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to systems for identification and characterisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-508123/54.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligo dT primer; gene expression analysis; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligo dT primer #2 used in method to study gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GENO-)
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BP; 20 A; 0 C; 1 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            45pp; English.
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RESULT 967
ADB04575/c
ID ADB045
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AC ADB045
XZ
DT 20-NOV
XX
DE Human
XX
KW Cytost
KW zinc f
                                                                                                                                                                                                                                                                                                                                                                                                                               CC of gene expression in one or more samples, comprising an identifier having a specific oligo-dT primer sequence, where the identimer comprises a CC detectable marker at its 5' end. The system is useful for identifier having a given in vivo or in vitro RNA sample, as well cas the relative differences in mRNA between 2 or more samples, where CC desired, for supporting discovery of new genes, and for identifying mRNAs that are expressed at different levels between 2 or more samples. The new compositions and systems that incorporate new strategies where more system or method addresses limitations of prior methods by comprising CC compositions and systems that incorporate new strategies where molecular or biochemical assay compositions and systems are linked to DNA or RNA compositions and systems and systems are linked to DNA or RNA compression. The system has the following advantages over existing compression. The system has the following advantages over existing composition in addition to information concerning changes or differences in formation in addition to information concerning changes or differences continued to the sequence sample for subsequent investigation by common molecular biology compositions and (d) does not require prior knowledge of the sequence of the organism under investigation and can be employed in concernion and can be employed in sequence represents an oligo of primer used in the method of the
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                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity Matches 17; Conserv
 Cytostatic; immunostimulant; gene therapy; vaccine; human; zinc finger protein; MDZ3; MDZ4; MDZ7; MDZ12; chromosome 7q22.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying and characterizing gene expression in samples, for identifying mRNAs expressed at different levels, comprises employing an identifier having a oligo-dT primer of a specific sequence and a detectable marker at its 5' end.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200236828-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligo
                                                         Human MDZ7 scanning oligonucleotide SEQ ID 5561.
                                                                                                 20-NOV-2003
                                                                                                                                       ADB04575
                                                                                                                                                                           ADB04575
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 24 BP; 3 A; 1 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to systems for identification and characterisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 11; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kane MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-2000; 2000US-0244933P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-2001; 2001WO-US045401.
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                                                                                                                                                                                                                                                                                               1246 TCTTTGTTTTGTTTTTTAATC 1266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     dT primer; gene expression analysis; primer; ss
                                                                                                                                                                                                                                                                           4
                                                                                                                                                                             standard; DNA; 25
                                                                                                                                                                                                                                                                           TTTTTTTTTTTTTTAAAC 24
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                                                                                                                                                                                                                                                                                                                                                   1.0%;
| llarity 81.0%;
| Conservative
                                                                                                 (first entry)
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Pred. No. 46
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                                                                                                                                                                                                                                                                                                                                                                                       Length 24;
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AANSO186/c
ID AANSO1
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AC AANSO1
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DT 25-MAR
DT 01-NOV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to novel human zinc finger-containing CC proteins and their coding sequences; MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is CC encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 6p21.3-22.2, CC MDZ7 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome (fight). The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, CC or in manufacturing a medicament for treating or preventing a disorder CC associated with decreased or increased expression or activity of MDZ3, CC MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic CC acids and proteins are also useful for diagnosing or monitoring a disease CC caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic CC acids can also be used as probes to detect and characterize gross CC alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are CC useful in constructing microarrays for measuring gene expression. The CC proteins are useful as therapeutic agents for gene therapy or as CC vaccines. The present sequence was used to illustrate the invention.
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chromosome 6p21.3-22.2; chromosome 16p11.2; chromosome 15q26.1; cancer;
developmental disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
01-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 25 BP; 4 A; 2 C; 3 G; 16 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-JUL-2002; 2002EP-00016874.
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   06-NOV-1984;
                                                                  03-JUL-1985
                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                              Immunoglobulin
                                                                                                                                                                                                                                                                                                              Sequence of human immunoglobulin gene enhancer region called unit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAN50186;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAN50186 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 8; SEQ ID NO 5561; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MDZ4, MDZ7 or MDZ12, e.g. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (AEOM-) AEOMICA INC
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                                                                                                                            EP146743-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                          (revised)
(first entry)
   84EP-00113371
                                                                                                                                                                                                                                                 gene; enhancer; promoter; expression cassette;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        81.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%;
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Pred. No. 3.
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RESULT 969
RAXK89795/c
ID AAXK897
XX AAX697
XX AAX697
XX Human
XX Human
XX Vaecul
KW KDR; h
KW tumour
KW fme-li
KW foetal
XX Homo s
XX Homo s
XX W09715
XX PF 25-OCT
XX 26-OCT
PR 11-JAN
XX (RIBO-PA (CHIR
XX Pavco
XX Pavco
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XX Nuclei
PT Nuclei
PT rheums
XX Claim
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Best Local S
Matches 15
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06-DEC-1983;
04-JUN-1984;
05-JUN-1984;
                                                                                                                                                                                                                                                                                                    KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; tems-like tyrosine kinase i; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The inventors claim a gene fragment comprising (a) the enhancer DNA segment and (b) a structural gene such as human D, V and J gene, and promoter. Using the gene fragments of the invention, transcription etranslation efficiency are enhanced. (Updated on 25-MAR-2003 to corr
                                       stability
                                                                         WPI; 1997-259017/23.
                                                                                                                                                          26-OCT-1995;
11-JAN-1996;
                                                                                                                                                                                             25-OCT-1996;
                                                                                                                                                                                                                     01-MAY-1997
                                                                                                                                                                                                                                             WO9715662-A2
                                                                                                                                                                                                                                                                                                                                            Vascular endothelial growth factor receptor; VEGF receptor; flt-1;
                                                                                                                                                                                                                                                                                                                                                                   Human flt1 VEGF receptor hammerhead ribozyme substrate #1090.
                                                                                                                                                                                                                                                                                                                                                                                            28-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX69795
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fragments derived from chromosomal DNA cuseful in transcription and translation
                         rheumatoid arthritis,
                                                                                                                                                                                                                                                                    Homo sapiens
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15; Conserv
                                                                                                                        RIBOZYME PHI
CHIRON CORP
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                                                 acid molecule modulating VEGF
                                                                                                                                                                                                                                                                                                                                                                                                                                            standard;
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                                                                                              Mcswiggen J,

    useful

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                                                                                                                                                                                                                                                                                            kinase 1; ss.
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83JP-00229037.
84JP-00113047.
84JP-00113834.
                                                                                                                                                          95US-0005974P.
96US-00584040.
                                                                                                                                                                                              96WO-US017480
                                                                                                                                  PHARM INC
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93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                            RNA;
                                    for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0 C; 0 G; 11 T; 0 U; 0 Other;
                        etc., in a human
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Pred. No. 6.6e+02;
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tumour angior
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                         patient
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                                    angiogenesis,
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                                                 gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                     psoriasis
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Claim 4;

Page 79; 218pp; English

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RESULT 970
AAA21311 E
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AC AAA21311;
XC AAA21311;
XC AAA21311;
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DT 19-JUN-200
XX Integrin a
XX Human; ary
XX integrin a
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Best Local S
Matches 15
The present invention describes enzymatic nucleic acid molecules with RN cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to AAA17167 and AAA17561 to AAA17622 represent ribozyme sequences for ARNT, and AAA17168 to AAA17560 and AAA17633 to AAA17684 represent their corresponding target sequences; AAA17685 to AAA18385 and AAA18386 to AAA19154 represent ribozyme sequences for Tie-2, and AAA18386 to AAA1908 and AAA19155 to AAA19222 represent their corresponding target sequences;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes nucleic acid molecules which modulate t synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VGGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase i (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; aryl hydrocarbon nuclear transport; ARNT; TIE-2; angiogenesis; integrin alpha 6 subunit; integrin subunit beta 3; hairpin ribozyme; hammerhead ribozyme; angiogenic factor; cytostatic; antidiabetic; ophthalmologic; antifilammatory; antiarthritic; antisoriatic; ARND; dermatological; RNA cleavage; cancer; diabetic retinopathy; arthritis; age related macular degeneration; inflammation; neovascular glaucoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               myopic degeneration; psoriasis; verruca vulgaris; angiofibroma; tuberous sclerosis; pot-wine stain; Sturge Weber syndrome; Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; ss
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Pred. No. 6.2e+02;
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Kippel-Trenaunay-Weber syndrome; Osler-Weber-Rendu syndrome; 8s.
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Pred. No. 6.2e+02;
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The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity, which specifically cleave RNA encoded by an aryl hydrocarbon nuclear transporter (ARNT) gene, an integrin subunit beta 3 gene, an integrin alpha 6 subunit gene, or a Tie-2 gene. AAA16775 to

Jarvis Parry T

, ' i

Beigelman L, M

Matulic-Adamic J, Reynolds M, Kisich K, Beigelman L, Mcswiggen JA, Karpeisky A, , Workman CT, Beaudry A, Sweedler D;

(RIBO-)

RIBOZYME PHARM INC.

Thompson J,

subunit gene,

Page 200; 305pp; English

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Matches 15
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09-JUN-1997;
03-JUL-1997;
22-AUG-1997;
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05-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; c-raf; A-raf; B-raf; hammerhead ribozyme; hairpin ribozyme; target; substrate; catalyst; modulation; expression; Raf gene; delivery; screening; identification; synthesis; deprotection; purification; cancer; screening; psoriasis; non-hepatic ascites; infection; genetic drift;
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No. 6.2e+02;
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ARESULT 973
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Screening for drugs that increase expression of the apolipoprotein A gene, which may then be useful for treating coronary artery disease.
                                                                                                                                                                                                                                                                                                                             29-SEP-1995;
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                                                                                                             WPI; 2000-038251/03
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RESULT 974
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC probes which comprise various different enhancer elements. These were CC used as competitor probes in competition gel mobility shift assays CC performed to determine the specificity of drug-inducible nuclear proteins CC for a fragment of the 5' flanking region of the human apolipoprotein AI (apo AI) gene comprising the sequence between bases -79 and -44 from the CC transcriptional start site (AAZ38357). This apo AI fragment contains two CC DRES (AAZ38355) and popposite orientations, and was used in the CC construction of plasmids pGL2 (apoAI-DRE)TK/luc (AAZ38352), which CC contains one copy of this fragment, and pGL2 (4xapoAI-DRE)TK/luc (AAZ38353) which contains four copies. The fragment (s) were inserted CC upstream of the HSV thymidine kinase (TK) promoter and luciferase CC reporter gene in the pGL2 TK/luc vector The pGL2 TK/luc vector also CC comprises a functional polyadenylation sequence downstream of the CC reporter gene. Plasmids pGL2 (apoAI-DRE)TK/luc and pGL2 (4xapoAI-DRE)TK/luc are used in a novel method for screening for a drug that CC increases expression of a gene for apolipoprotein (apo AI), which is CC associated with coronary artery disease. Such plasmids may be transformed CC untreated cells. The method is useful for screening and identifying drugs that increase apo AI gene expression, the identified drugs would be cuseful for treating coronary artery disease. The method is simple, rapid, candular contains coronary artery disease. The method is simple, rapid, candular contains form and coronary artery disease. The method is simple, rapid, candular contains coronary artery disease. The method is simple, rapid, candular coronary artery disease. The method is simple, rapid, candular coronary artery disease. The method is simple, rapid, candular coronary artery disease. The method is simple, rapid, candular coronary artery disease.
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Best Local
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Reynolds
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                                                                                                                                                                                                                                                                                                     20-APR-1998;
23-JUN-1998;
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                               WPI; 2000-013248/01
                                                                                                 Matulic-Adamic
                                                                                                                                                                                                                                  (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                      19-APR-1999;
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                                                                                                                             Beigelman L, Mcswiggen JA,
Zwick M, Jarvis T, Woolf T,
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98US-00103636.
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Haeberli
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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                     Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease;
                     New nucleic acids that
                                                          WPI; 2000-013248/01.
                                                                                                             Thompson JD, Beigelman L, Mcswiggen JA, Reynolds M, Zwick M, Jarvis T, Woolf T,
                                                                                                                                                                                                          20-APR-1998;
23-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                   anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oestrogen receptor hammerhead ribozyme target sequence SEQ ID NO:1944.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acids that interact, and optionally cleave, target sequences,
                                                                                             Matulic-Adamic J;
                                                                                                                                                                       (RIBO-) RIBOZYME PHARM INC
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15; Conserv
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98US-00103636.
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                     interact,
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Pred. No. 6.2e
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                     and optionally cleave, target sequences,
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. 6.2e+02;
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Haeberli
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC gene, are used to treat cancer (particularly of breast or endometrium), cc in vivo or by transforming cells ex vivo and implanting treated cells, or cc for other conditions associated with levels of oestrogen receptor. CC Because of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with alterations in phenotype, comparticularly for identification of therapeutic targets, and as research cc used with DNA). The combination of modifications in (A) improves cc used with DNA). The combination of modifications in (A) improves cc eaststance to nucleases, binding affinity and/or activity. AAA23503 to CAAA24747 represent oestrogen receptor hammerhead ribozyme sequences, and cc AAA24748 to AAA2592 represent their corresponding target sequences, and CC AAA24748 to AAA25105 represent toestrogen receptor hairpin ribozyme sequences. AAA25219 to AAA26211 represent their corresponding target sequences and cc sequences. AAA26210 to AAA26271 represent their corresponding target sequences and cantisense oligonucleotides used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                             Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor prointerferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes nucleic acids (A) that interact stably with a target sequence and contain at least one phosphoro(di)thicate link, having endomuclease activity. (A), and more generally any catalytic nucleic acid (A') that modulates expression of the oestrogen receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17 BP; 1 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
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                                                                                                                                                                    WPI; 2000-647423/62.
                                                                                                                                                                                                    Blatt L, Zwick M,
                                                                                                                                                                                                                                                                       12-APR-1999;
                                                                                                                                                                                                                                                                                                         11-APR-2000; 2000WO-US009721
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                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ribozyme; erythropoietin; granulocyte colony stimulating factor;
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                                                               Claim 42; Page 123; 164pp; English.
                                                                                                                                                                                                                                       (RIBO-)
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                                                                                                                                                                                                                                       RIBOZYME PHARM INC.
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Pred. No. 6.2e+02;
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The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcripti

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RESULT 977
AAFO6140
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XX AAFO61
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XX Homer
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KW Ribozy
KW Ribozy
KW Interf
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XX W22000
XX Homo 8
XX W11-APF
XX H1-APF
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XX H1-APF
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Best Local
                                                                                                                                                                                                                                                                                 The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of pressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor production alpha and erythropoletin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-647423/62.
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                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 42; Page 123; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-APR-1999;
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                                                                                                                             Score 14.4;
Pred. No. 6.
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Pred. No. 6.
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                                                                                                                                                      DB 1; Length 17;
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RESULT 978

Sequence 17

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                                                                          CC protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 CC lcan be used in gene therapy and vaccine production. The hGDMLP-1 cc nucleic acids in samples, as amplification substrates and quantify cc hGDMLP-1 nucleic acids in samples, as amplification substrates, to c provide initial substrates for the recombinant engineering of hGDMLP-1 cc protein variants having desired phenotypic improvements, and for c expressing the proteins. The hGDMLP-1 proteins or polypeptides may be cused as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration cand/or amount specifically of hGDMLP-1 proteins, as specific blomolecule capture probes for surface-enhanced laser description ionisation, as construction, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The colorated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence in the expression of the present invention. N.B. chapter as content on the present invention. N.B. the sequence data for this patent did not form part of the printed capture in the many obtained in electronic format directly from MIPO care from wino intronic handled in electronic format directly from MIPO.
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30-JAN-2001;
30-JAN-2001;
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30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAY-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human GDMLP-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (AEOM-) AEOMICA INC.
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21-SEP-2000;
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                                                    ftp.wipo.int/pub/published_pct_sequence
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                  The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynuclectide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as
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as specific biomolecule capture probes for surface-enhanced laser
sorption ionization, comprises human myosin-like protein hGDMLP-1.
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Pred. No. 6.2e+02;
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RESULT 980
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AC ABNO22
XX ABNO22
XX 29-MAY
XX Human;
XW Human;
XW muscle
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XX 26-MAY
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30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US0006667.
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21-SEP-2000;
27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                         New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                            Disclosure; SEQ
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                                                                         ID NO 2271; 214pp;
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Pred. No. 6
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                                                                            English.
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The present invention protein 1 (hGDMLP-1).

invention describes a human genome-derived myosin-like MGDMLP-1). The protein and polynucleotide sequences of

hGDMLP-

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RESULT 981
ABNO22
XX ABNO22
XX ABNO22
XX ABNO22
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XX ABNO22
XX Human
XX Human
XX Homo 8
XX WO2001
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PR 27-SEP
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Best Local :
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21-SEP-2000;
27-SEP-2000;
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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                                                          (AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                        30-JAN-2001;
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2001WO-US000665.
2001WO-US000667.
2001WO-US000667.
2001WO-US000668.
2001WO-US000669.
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2001WO-US000661.
2001WO-US000662.
2001WO-US000663.
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2000US-0234687P.
2000US-0236359P.
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   Penn SG,
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   Chen
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   Shannon
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                  therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                    Sequence 17
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                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                        ftp.wipo.int/pub/published_pct_sequence
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                                                                   396 GCCGAAGGCCCGCAGG 411
17 GCCGAAGGCCCGGAGG
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15; Conservat
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Pred. No. 6.2e+02
0; Mismatches
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RESULT 982
ACN12480
ID ACN124
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ACN124
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ACN122
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ACN122
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ACN122
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WNV mi
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WNV; W
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                                                                                                       (RIBO-) RIBOZYME PHARM INC (BLAT/) BLATT L.
                                                                                                                                                                                                                                                                                                                                                                 19-OCT-2001; 2001WO-US048350.
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Best Local
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New nucleic acid molecule that modulates replication of West Nile (WNV), useful for treating a condition related to WNV infection e. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                                                                                                                           (RIBO-)
(BLAT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; lifection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                         WPI; 2002-706994/76.
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BLATT L.
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Pred. No. 6.2e+02;
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RESULT 984
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephallitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                      New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
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                                                                                                                                                                                              Blatt
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                                                                                                                                                                WPI; 2002-706994/76.
                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC. (BLAT/) BLATT L. (MCSW/) MCSWIGGEN J A.
                                                                                                                                                                                                                                                                                                                 19-OCT-2001; 2001WO-US048350
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                                                                       1199; 495pp;
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Pred. No. 6.2e+02
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The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (MNV). The nucleic acid molecules are useful for treating a condition related to MNV infection e.g. pancreatitis, encephalitis, myocarditis, meningitis, neurologic infection, hepatitis,

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The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, encephalitis, myocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic molety. Nucleic acid molecules SEQ ID NO 1 to 37080
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                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; myocarditis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-OCT-2001; 2001WO-US048350
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                         The invention relates to nucleic acid molecules that modulate replication of the West Nile Virus (MAV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, encephalitis, mocarditis, meningitis, neurologic infection, hepatitis, niver failure, hepatocellular carcinoma or cirrhosss. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least three of the 5' terminal nucleotides and a 3' end modification of a 3'-3' inverted abasic molety. Nucleic acid molecules SEQ ID NO 1194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid molecule of the invention
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(BLAT/)
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                                                                                                                                                     of the West Nile Virus (WNV). The nucleic acid molecules are useful for treating a condition related to WNV infection e.g. pancreatitis, encephalitis, myocarditis, meningitis, neurologic infection, hepatitis, liver failure, hepatocellular carcinoma or cirrhosis. The nucleic acid molecule is selected from the group of ribozymes consisting of Hammerhead, Inozyme, G-cleaver, DNAzyme, Amberzyme and Zinzyme. The nucleic acid molecules further comprise at least five ribose residues, at least three of the 5' cerminal nucleotides and a 3' end modification of a 3' 3' inverted abasic moiety. Nucleic acid molecules SEQ ID NO 1 to 37080 are claimed; however, SEQ ID NO 2194-2206 and 17502-17514 are not given in the specification. The present sequence is that of a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acid molecule that modulates replication of West Nile Virus (WNV), useful for treating a condition related to WNV infection e.g. pancreatitis, meningitis, hepatocellular carcinoma or cirrhosis.
                                                                                                                Sequence 17
                                                                                                                                                                                                                                                                                                                                                                  Claim 23; SEQ ID NO 7862; 495pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-706994/76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-OCT-2000; 2000US-0242411P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          West Nile Virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WNV; West Nile Virus; antiinflammatory; cytostatic; hepatotropic; virucide; neuroprotective; antibacterial; replication; pancreatitis; encephalitis; meningitis; infection; hepatitis; liver failure; cancer; cirrhosis; Hammerhead; Inozyme; DNAzyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACN07859 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                       The invention relates to nucleic acid molecules that modulate replication
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-OCT-2001; 2001WO-US048350
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC.
(BLAT/) BLATT L.
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                                                                       Similarity
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                                                                                                                                              of the invention
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                            CGCTGGTGAATGGAGT 711
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                                                                                                                 BP; 4
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                                                         Conservative
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                                                                       1.0%;
                                                                                                                 7 C; 3 G; 0 T; 3 U; 0 Other;
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                                                                     Score 14.4; DB 1;
Pred. No. 6.2e+02;
                                                           Mismatches
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                                                                                   DB 1;
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RESULT 988
ABT36080/c
                                                                     The invention relates to a novel isolated 17 mer nucleic acid sequence, cc given in the specification, a sequence containing at least 15 consecutive contices in the specification, a sequence with, after optimal cc alignment, at least 80 % identity to the 17 mer sequence that cc hybridizes to them under highly stringent conditions, or the complement cof any of them, or the corresponding RNA. The novel isolated mucleic acids of the invention are useful as probes and primers for detecting, cc identifying, quantifying and/or amplifying a nucleic acid, e.g. as one cc component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, cc polypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for c preparation of pharmaceuticals for prevention and/or treatment of viral cd diseases that are characterised by development of tumours or cell cd degeneration, specifically cancer but also Albeimer's disease and comparation procession of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these chipseptide and antibodies are useful as components of these chipse. The nucleic acid sequences of the invention can be used in gene chips. The nucleic acid sequences of the invention can be used in gene characterise and contained in the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene chipse. The nucleic acid sequences of the invention can be used in gene characterise and contained in gene characterise and contained in gene contained the invention can be used in gene contained the invention can be used in gene contained the invention.
Query Match
Best Local S
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                                                                 Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 233; 720pp; French.
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                                                                   BP; 6 A; 3 C; 4 G; 4 T; 0 U; 0 Other;
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  1.0%;
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                         Score 14.4; DB 1;
                         Length 17;
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16

1315 TTTTCAGAGACAGATC 1330

Similarity

0 Pred.

Mismatches No.

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Gaps

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6.2e+02

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ABT35552
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Cytc
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RESULT 990
ABZ64551/c
ID ABZ645
                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel isolated 17 mer nucleic acid sequence, given in the specification, a sequence containing at least 15 consecutive CC nucleotides from the 17 mer sequence with, after optimal CC alignment, at least 80 % identity to the 17 mer sequence that CC hybridizes to them under highly stringent conditions, or the complement CC soids of the invention are useful as probes and primers for detecting, CC identifying, quantifying and/or amplifying a nucleic acid, e.g. as one CC component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, colypeptides, vectors containing the nucleic acids, cells containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaccuticals for prevention and/or treatment of viral CC diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and cc diseases. The polypeptides can also be used to generate antibodies, and cohypeptide and antibodies are useful as components of these caid sequences of the invention can be used in gene cherapy. This polynucleotide sequence represents a tumour suppression cc related human fukutin oligonucleotide of the invention can be used in gene
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Best Local Similarity
Matches 15; Conserv
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   ABZ64551 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-SEP-2001; 2001FR-00011978
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-JUN-2003 (first entry)
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                                                                                                                                                                                                                   ATTITATTTCCCTT 1363
                                                                                                                                                            ATCTTATTTCCCTT 17
                                                                                                                                                                                                                                                                                                                                                                                                            BP; 2
                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                         A; 4 C; 1 G; 10 T; 0 U; 0 Other;
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Pred. No. 6.2e+
0; Mismatches
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RESULT 991
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel short interfering RNA (siRNA) nucleic acid molecule or an enzymatic nucleic acid molecule, that modulates expression of a nucleic acid molecule encoding HERZ, K-Ras, H-Ras, N-Ras, human immunodeficiency virus (HIV) or a component of HIV. The nucleic acid molecule of the invention has cytostatic, anti-HIV, and anti-rheumatic activity. The nucleic acid molecules are useful for reducing HER2, K-Ras, H-Ras, and HIV activity in a cell. The nucleic acids are also useful for treating breast, ovarian, colorectal, lung, prostate, bladder, or panneatic cancer, and HIV infection, and AIDS. The sequences shown in ABZ59889 - ABZ6216, ABZ64544 - ABZ65531, ABZ66520 - ABZ66524, ABZ66530 - ABZ66585 represent substrate/target sequences for the human relations of the inventor of the substrate of target sequences for the human relations.
Nucleic acid molecule; Hepatitis C virus; HCV; Hepatitis B virus; HBV; RNA stability; RNA expression; RNA synthesis, antisense; enzymatic nucleic acid; hammerhead ribozyme; DNazyme; inozyme; zinzyme amberzyme; G-cleaver ribozyme; decoy molecule; aptamer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; ribozyme; short interfering RNA; siRNA; HER2; K-Ras; enzymatic nucleic acid; H-Ras; N-Ras; HIV; cytostatic; anti-HIV; anti-rheumatic; cancer; AIDS; ss.
                                                                                 HCV DNAzyme
                                                                                                                24-SEP-2003
                                                                                                                                                                               ACD60861 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel short interfering RNA and enzymatic nucleic acid useful for treating cancer, modulates the expression of a nucleic acid encoding HER2, K-Ras, H-Ras, N-Ras, and human deficiency virus sequences.
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06-JUN-2001; 2001US-0296249P
10-SEP-2001; 2001US-0318471P
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                                                                                                                                                ACD60861;
                                                                                                                                                                                                                                                                                                                                                                                                                              ribozymes of the
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                                                                                                                                                                                                                                                                                               GGGGCGCGCCTGAC 484
                                                                               substrate sequence #2047.
                                                                                                                                                                                                                                                                                                                                                                                            BP; 1 A; 9 C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                invention
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                                                                                                                                                                                                                                                                                                                                             Score 14.4; DB 1;
Pred. No. 6.2e+02;
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RESULT 992
ADB44694
ID ADB44694
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                                                                                                                                                                                    The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, C inozymes, zinzymes, and G-cleaver ribozymes. Also disclosed are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV C DNA. The nucleic acids may be used to modulate the expression of HBV compounds and/or potential therapies directed against HBV, and compounds compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HCV broader or minus strand DNAzyme sequences disclosed in the present
                                                                                                                            Query Match
Best Local S
Matches 15
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08-JUN-2001;
08-JUN-2001;
24-OCT-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HBV reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic;
                                                                                                                                                                                   Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 270; 387pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel compound useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-229207/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Blatt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-MAR-2002; 2002WO-US009187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200281494-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     virucide; antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hepatocellular carcinoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Draper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MORR/)
(PAVC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RIBO-)
(BLAT/)
(MACE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LEEP/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MCSW/)
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                                                                                                  961
                                                                        17
                                                                                                                            15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DRAPER K.
ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RIBOZYME
BLATT L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LEE P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PAVCO P.
                                                                                                  TCGCGCGGCCCCGGCG 976
                                                                      TCGCGCGGCACCGGCG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Macejak D,
Roberts E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; 2001US-00817879.
; 2001US-00877478.
; 2001US-0296876P.
; 2001US-0335059P.
                                                                                                                                                                                   BP; 1 A;
                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2001US-0337055P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PHARM INC
                                                                                                                                          1.0%;
                                                                                                                                                                                   7 C; 7 G; 0 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for treating cirrhosis, liver failure, oma, or condition associated with hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mcswiggen
                                                                        N
                                                                                                                            0
                                                                                                                                        Score 14.4; DB 1
Pred. No. 6.2e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               J, Morrissey D,
                                                                                                                              Mismatches
                                                                                                                                                                                   2 U; 0 Other;
                                                                                                                                                     DB 1;
                                                                                                                                                       Length 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pavco
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                                                                                                                            Gaps
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standard;

DNA; 17

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27-JUN-2003 ACC54422;

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The invention relates to the isolation of 6327 nucleotide sequences, and consecutive nucleotides of these nucleotides, a sequence having at least 80% identity, after optimal alignment, with the nucleotides, a sequence that hybridizes under stringent conditions with the nucleotides, or the complement, or corresponding RNA, of the nucleotides. The nucleotides are used as probes or primers for detecting, containing the nucleotides are used as probes or primers for detecting, containing the vectors of nucleotides involved in tumour suppression or reversion, apoptosis and or viral resistance, to produce recombinant polypeptides, and to prepare transgenic animals, as experimental models. The nucleotides (also vectors containing the mand containing the vectors), the encoded polypeptides and antibodies (ab) against the polypeptide are useful for prevention and/or treatment of viral infections or diseases characterized by development of tumours corell degeneration (e.g. Alzheimer's disease or schizophrenia).

Chalysis of the expression of the nucleotides can be used for diagnosis and/or prognosis of these diseases. The nucleotides and polypeptides can potentially useful for treating diseases associated with abnormal corporation of the nucleotides.
                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                            Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New nucleic acid encoding human prostate membrane-specific antigen, useful e.g. for treatment of tumors and viral infection, also related polypeptide and antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic; antiviral; neuroprotective; nootropic; neuroleptic; ss;
primer; probe; tumour suppression; tumour reversion; apoptosis;
virus resistance; transgenic animals; Alzheimer's disease; schizophrenia;
ACC54422
                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 2 A; 4 C; 1 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 618; 771pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Telerman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-SEP-2001; 2001FR-00011981
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tumour suppression/reversion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-SEP-2002; 2002WO-IB004219
                                                                                                                                                                                                      1348
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                                                                                                                                                                                                                                                                                            Similarity
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standard;
                                                                                                                                                                                       ATTTTTATTTTCCCTT
                                                                                                                                                  ATCITIATTITCCCTT 17
                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amson R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
DNA; 17
                                                                                                                                                                                                                                                                                            1.0%;
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   ₽P
                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                         Score 14.4; DB 1;
Pred. No. 6.2e+02;
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                                                                                                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                              1.
                                                                                                                                                                                                                                                                                                                     Length 17;
                                                                                                                                                                                                                                                                    Indels
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RESULT 994
AD184801/C
ID AD1848
XX AD1848
XX AD1848
XX O3-JUN
XX S8; en
KW HCV IN
XX HEQUIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                       HCV DNAzyme substrate sequence #2047.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADI84801;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence represents an isolated nucleic acid sequence associated with tumour suppression or regression, apoptosis or virus resistance. Invention relates to these sequences or sequences having at least 80% identity to them, and polypeptides encoded by the sequences or polypeptides having 80% identity to the polypeptide sequences. The invention is used to diagnose or treat viral disease or disease characterized by development of tumour cells or cellular degeneration
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New nucleic acid sequences associated with tumor suppression, regression, apoptosis or virus resistance are useful to diagnose and treat viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-JUN-2001; 2001FR-00008139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tumour regression; apoptosis; virus cellular degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADI84801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FR2826373-A1
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  (BLAT/)
                                                                               18-DEC-2000;
                                                                                                                                                                                                                                           US2003125270-A1
                                                                                                                                                                                                                                                                                         Hepatitis C virus.
                                                                                                                                   18-DEC-2000; 2000US-00740332.
                                                                                                                                                                                       03-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MOLE-) MOLECULAR ENGINES LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tumour suppressor; antitumour; cytostatic; tumour suppression; our regression; apoptosis; virus resistance; diagnosis;
                                                                                                                                                                                                                                                                                                                                          enzymatic nucleic acid; RNA cleavage; hepatitis infection; type I interferon; DNAzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1348
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l Similarity
15; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard; RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTTTATTTTCCCTT 1363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  development of tumor cells and cell degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATCTTTATTTTCCCTT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%;
illarity 93.8%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 2 A; 4 C; 1 G; 10 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Telerman
                                                                               2000US-00740332
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 14.4; DB 1;
Pred. No. 6.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 17;
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                                                                                                                                                                                                                                                                                                                                                                           virus;
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RESULT 995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to an enzymatic nucleic acid molecule which specifically cleaves RNA derived from hepatitis C virus (HCV), in which the binding arms of the enzymatic nucleic acid molecule comprises sequences complementary to any of the defined substrate sequences given in the specification. The nucleic acid molecule may be administered for the treatment of HCV infections, especially in combination with type I interferons. The present sequence represents a HCV DNAzyme substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Enzymatic nucleic acid molecules which specifically cleave from hepatitis C virus (HCV), useful for the treatment of H especially in combination with type I interferon therapy.
The present sequence is one of a large number of antisense oligonucleotides which is directed against one of four human receptor RNA activator (SRA) nucleic acid sequences. Two seri
                                                                                                                                                                              (ISIS-)
                                                                         Antisense compounds targeted to steroid receptor RNA activator useful diagnosis, prophylaxis and treatment of diseases associated with the steroid activator, such as infection, inflammation or tumor formation
                                                                                                                                                                                                                                                                                              US6107092-A
                                                                                                                                                                                                                                                                                                                                                             Human;
                                                                                                                                                                                                                                                                                                                                                                                     Antisense oligonucleotide ISIS# 30213
                                                                                                                                                                                                                                                                                                                                                                                                              04-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 2047; 198pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ROBE/) ROBERTS E. (PAVC/) PAVCO P A. (MACE/) MACEJACK D.
                                                                                                                            WPI; 2000-586211/55
                                                                                                                                                      Cowsert
                                                                                                                                                                                                                    29-MAR-1999;
                                                                                                                                                                                                                                            29-MAR-1999;
                                                                                                                                                                                                                                                                     22-AUG-2000
                                                                                                                                                                                                                                                                                                                                                SRA inhibitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA92546;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17
                                                                                                                                                                                                                                                                                                                                              SRA; steroid receptor RNA activator; cytostatic; antiinflammatory; hibitor; cancer; infection; antisense oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
15; Conserv
                                                                                                                                                                              ISIS PH
BAYLOR
                                                                                                                                                      <u>`</u>
                                                  Co1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TCGCGCGGCACCGGCG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 1 A; 7 C; 7 G; 0 T; 2 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                              PHARM INC.
OR COLLEGE MEDICINE
                                                 41; 47pp; English
                                                                                                                                                      Bennett
                                                                                                                                                                                                                                                                                                                                                                                                              (first entry
                                                                                                                                                                                                                    99US-00280409
                                                                                                                                                                                                                                            9905-00280409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
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                                                                                                                                                      O'malley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.4; DB 1;
Pred. No. 6.2e+02;
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                                The present invention describes a set of peroxidase genes found in plants, especially rice, and their homologues, modified forms and fragments, where the sequences of the peroxidase genes in the set are given in AAH44071 to AAH44091. Also described are: (1) promoters for the control of the gene set; (2) the preparation of cassette vectors using the genes and promoters; (3) analysis of plant characteristics using the peroxidase set by isolating RNA from the plant, binding the RNA to a membrane, mixing with a labelled peroxidase gene set, incubating, and detecting the label signal to show which genes in the set are expressed in the sample plant; and (4) DNA microarrays for peroxidase gene expression analysis. The set of genes are used for the analysis of the pattern of peroxidase gene expression in particular rice plants and their component tissues and under different environmental conditions, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ohashi Y,
Hiraga S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -30 oligodeoxynucleotides with a phosphorothioate backbone. The second series comprised chimeric oligonucleotides composed of a central gap region, consisting of ten 2'-deoxynucleotides, which was flanked on both sides by four-nucleotide wings. The wings were composed of 2'-methoxyethyl (2'-MOE) nucleotides. Both series contained the same nucleotide sequences. The antisense compounds are useful for research, diagnosis, treatment and prophylaxis to prevent or delay infection, inflammation or tumour formation. Therapeutically the oligonucleotides are highly safe and are effectively administered to humans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oryza sativa; rice; peroxidase; POX; characteristic; gene expression; modification; plant; bacterial infection; Magnaporthe grisea; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; Page 62; 258pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rice under varying
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NORQ ) JAPAN MIN AGRIC FORESTRY & FISHERIES
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Pred. No. 5.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      for analysis of peroxidase expression in production of rice plants with desired
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modification of rice plants to provide desired specificities

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ABS52682/
ID ABS5
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       CC system comprising translation factors and transfer ribonucleic acidd (C (tRNA) species which translate exogenously added messenger RNA (mRNA) CC with highly selective incorporation at each codon to form a peptide or a peptidominetic product when the system includes one or more tRNA species CC charged with a synthetic amino acid or amino acid analogue. The CC charged with a synthetic amino acid or amino acid analogue. The CC peptide or protein ligands or catalysts, such as insulin, growth hormone CC peptide or protein, and for pure ribosome display and pure mRNA display CC selection experiments. The translation process provides a simplified, CC highly purified system that offers potentially improved routes to all CC peptides and proteins currently synthesised by alternative routes. This CC maintaining purified components and trace contaminants or insefficient CC processivity. There are several advantages associated with performing CC post-translational modification of peptides, lack of proteases which CC ontaminants in the selection steps. The present sequence represents a contaminants in the selection steps. The present sequence represents a contaminants in the selection steps. The present sequence represents a contaminants of in the selection steps.
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                   e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                    A new reconstituted cell-free translation system comprising translation factors and tRNA species capable of translating exogenously added mRNAs useful for the synthesis of peptides or protein ligands or catalysts,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      peroxidase gene expression to impart particular characteristics to the plants such as response to bacterial infection by Magnaporthe grisea. present sequence represents a PCR primer/probe for a rice peroxidase (POX) gene, which is used in an example from the present invention
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                                                                                                                                                                                                                                                                                                                                               This invention relates to a novel reconstituted cell-free translation
                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 15; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Forster AC,
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oligonucleotide used
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Pred. No. 5.9e+02;
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                                                          CC contacting a target nucleic acid having one or more sequences selected from 5 3581 base pair sequences in a biological sample with at least one creagent or a series of reagents. The method is useful for detecting the methylation state of the 5 and promoter region of the gene DD3 within a collection of the series of the 5 and promoter region of the gene DD3 within a collection of the gene DD3 within a complementary sequences. The set of oligomers is also useful for complementary sequences. The set of oligomers is also useful for collecting the methylation state and/or collections within SEQ ID NO. 1 and its complementary sequences. The set of all cpd dinucleotides within SEQ ID CC amplification of DNA sequences selected from SEQ ID NO. 1 and its complementary sequences selected from SEQ ID NO. 1 and its complementary sequences selected from SEQ ID NO. 1 and its complementary sequences selected from SEQ ID NO. 1 and its complementary sequences with the methylation state of the gene DD3 comprising diseases associated with the methylation state of the gene DD3 comprising at least one nucleic acid. The methods, nucleic acids, colligonucleotide or PNA-oligomer, kit, array or the set of coll proliferative disorders. The present collison be used for the characterization, classification. The collipse of cell proliferative disorders. The present collison collection oligonucleotide of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detecting methylation of 5' and promoter region of DD3 gene for diagnosing proliferative disorders comprising contacting target acid with a reagent that distinguishes between methylated and methylated CpG dinucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to detecting the methylation state of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 6; SEQ ID NO 76; 56pp; English.
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Pred. No. 5.9e+02;
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                          CC Incus AST-1 for asthma and other IgE mediated diseases mapped to the CC human chromosome 7p14-p15. Specifically, it refers to two overlapping CC genes namely GPRA (G-protein coupled receptor for asthma susceptibility) CC and AAA1 (asthma associated alternatively spliced gene 1). The present CC invention describes identifying single nucleotide polymorphisms, as well as insertion or deletion polymorphisms, occurring at different positions in the AST-1 locus, and furthermore providing vectors, host cells, CC primers and provides a kit to diagnose or assess predisposition to CC Accordingly, it provides a kit to diagnose or assess predisposition to CC asthma, chronic obstructive pulmonary disease or cancer and other IgE mediated diseases including rhinitis and dermatitis, such that derived CC pharmaceutical compositions exhibit cytostatic and antiasthmatic CC activities. Furthermore, it provides a transgenic animal comprising the asthma locus-1 (AST-1) DNA. This oligonucleotide sequence is a 5' splice junction of the human AAA1 gene, given in Table 11 of the invention.
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03-JAN-2003; 2003US-043789SP.
26-MAR-2003; 2003US-0458767P.
09-JUL-2003; 2003US-0486000P.
                                                                                                                                                                                                                                                                                                                            New GPRA polypeptides, useful in preparing a composition for diagnosing, treating or preventing asthma, other IgE-mediated disease, chronic obstructive pulmonary disease or cancer.
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                                                                                                                                                                                                                                                                                               7; Page 83; 265pp; English.
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Pred. No. 5.9e+02;
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                                                                                                                                                                                                                                                                  New hooked or semi-hooked oligonucleotides (see AAQ68869-71, AAQ68873, AAQ68875, AAQ68877, AAQ68879 and AAQ68880) are useful as therapeutic antisense molecules for treating dermatological disorders (e.g. atopical dermatitis, psoriasis, melanoma, T cell lymphoma etc.) viral infections (e.g. herpes simplex, papilloma, hepatitis or HIV); or cancer (when directed against an oncogene), due to their ability to hybridise with target nucleic acid. They can be used ex vivo, e.g., to treat bone marrow grafts. They can also be used for diagnosis or in cosmetics e.g. to block mRNA coding proteins involved in the ageing process such as collagenase or elastase. This linear antisense oligonucleotide is used as a control to see whether the hooked and semi-hooked oligonucleotide sexhibit a greater resistance to exonucleases than linear oligonucleotides. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct PN field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New oligo:nucleotide(s) self paired at one or both ends - have improved resistance to nuclease(s) and reduced toxicity, useful as anti:sense molecules for treating dermatological disorders, virus infections,
                                                                                                                                                                                                                 Sequence 20
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                                                                                                        Score 14.4; D
Pred. No. 5.3e
0; Mismatches
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                                                                                                                                                                                                                    G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     control in antisense therapy.
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                                                                                                                               5.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Poddevin
                                                                                                                                                              DB 1;
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                                                                                                                                                              Length 20;
                                                                                                              Indels
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RESULT 1001
AAZ03168/c
ID AAZ0316
XX AAZ0316
XX AAZ0316
XX AAZ0316
XX AAZ0316
XX Vaccin
DT 07-OCT-
DE PCR pri
XX Vaccin
KW Vaccin
KW PARATTC
XX NONgor
KW DARTTC
CS Synth
COS Chlam
XX NO992
XX NO992
XX NO992
XX I0-JU
XX 28-N
PR 17-D
PR 17-
AAX97150
ID AAX9
XX
AC AAX9
XC
DT 13-9
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Resi
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Best Local
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17-DEC-1997;
04-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFs) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFs encode polypeptides (see AAY36754-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, pendymitis, cervicitis, salpingitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogranulomatosis. The polypeptides of the invention may be of use in treating these
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis; nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer; bartholinitis; pneumopathy; venereal lymphogranulomatosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
Chlamydia trachomatis.
                                                       PCR primer used to amplify an ORF of Chlamydia pneumoniae.
                                                                                                                                                                          AAX97150;
                                                                                                                                                                                                                               AAX97150 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 1584; 1755pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genome sequence of Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-371125/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer used
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                                                                                                                  13-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9928475-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                     1182 TGACGCCGATTCGTGT 1197
                                                                                                                                                                                                                                                                                                                                                                            19 ÍGÁTGCCGÁTTCGTGT 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                               (first
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97FR-00016034.
98US-0107077P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; 6 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                               entry)
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Pred. No. 5.3e-
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4; DB 1;
5.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
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Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis;

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RESULT 1003
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Best Local
                                                                                                                                              Human; DNA methyltransferase; DNA MeTase; antisense oligonucleotide; modulation; inhibition; gene expression; combination therapy; p16; histone deacetylase; HDAC; thymidylate synthase; tumour suppressor; methylation; gene therapy; tumour; cytostatic; antiasthmatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAY34584-AAY35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae nucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising
                                                                                                                                                                                                                 Human histone deacetylase HD2 antisense oligonucleotide SEQ ID NO:51:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX91991-X97517 represent PCR primers used to amplify open reading frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX91990). C. pneumoniae causes respiratory disease such as
  19-OCT-1998;
                                                                                                                                                                                                                                                                      AAA55806;
                                                                                                                                                                                                                                                                                               AAA55806 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genome sequence of Chlamydia pneumoniae
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04-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine; neutralising epitope; PCR primer; ss.
                           19-OCT-1999;
                                                                                WO200023112-A1
                                                                                                         Homo sapiens
                                                                                                                                    antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   epitope of C. pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 1881; Disclosure; 1912pp; English
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                                                                                                                                                                                                                                                                                                                                                                                    TTGCTGTTGCTGCTGC 740
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pneumoniae
  98US-0104804P
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98US-0107078P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98WO-IB001890
                           99WO-US024278
                                                                                                                                    inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                1.0%;
                                                                                                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 14.4;
Pred. No. 5.
                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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21-APR-1999;

99WO-IB000822.

28-OCT-1999. WO9954500-A2 Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; dentification; characterisation; haplotyping; hybridisation; dentification; SNP; PCR primer; amplification; single nucleotide polymorphism; SNP; PCR primer;

Human biallelic marker downstream amplification primer

SEQ ID

NO:9989

0

(first entry)

AAZ75633 standard; DNA; 20

diagnosis; ss.

sapiens

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AAZ7563
ID AAZ7563
XX AAZ7563
XX ID -SEP-
XX ID-SEP-
XX Human b
XX Human p
KW Human genomic
KW haploty
KW haploty
KW haploty
KW diagnos
XX Homo sa
XX Homo sa
XX Homo sa
XX Y SEP-
XX X Z SEP-
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CC inhibition of the gene. These may also be used to activate silenced genes to provide missing gene functions and improve a given condition.

CC purthermore, the methods and compositions are useful as probes of the composition. Anassystem; and to evaluate the effect of inhibiting gene activity or expression. Anassystem to Anassystem the effect of inhibiting gene activity or expression. Anassystem to analysis of the present oligonucleotide sequences which are used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 0 A; 7 C; 7 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TGCTGCTGCTGCC 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Macleod AR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.4; DB 1;
Pred. No. 5.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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RESULT 1005
ABL57552
ID ABL5755
XX ABL5755
XX ABL5755
XX Synthet
XX Concent
KW Concent
KW Polymex
XX Synthet
XX Synthet
XX COCT-
YN EP10467
XX Synthet
XX 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CAA265654 to AA269578 represent human biallelic markers from the present CC invention, which contain a polymorphic base at position 24 of their CC nucleotide sequences. AA269579 to AA277440 represent amplification primers for the biallelic markers. The biallelic markers of the invention CC have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies CC which are useful in determining the genetic basis for disease states. CC compositions and methods of the invention can also be useful for the CC identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the CC ifferential efficacious responses to and side effects from CC unsease actual agents acting on a disease as well as other treatment.

CC N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and CC agents invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               concentration; quantification; mutation detection; polymorphic; polymerase chain reaction; PCR: RR
                            WPI; 2000-657765/64
                                                                                                                                                                                                                                                      20-APR-1999;
                                                                                                                                                                                                                                                                                                  20-APR-2000; 2000EP-00108643
                                                                                                                                                                                                                                                                                                                                                        25-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                       EP1046717-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic deoxyribooligonucleotide poly s.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABL57552;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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23-NOV-1998;
                                                                                                                                                 (NIBI-) JAPAN BIOINDUSTRY ASSOC
(AGEN ) AGENCY OF IND SCI & TECI
(KANK-) KANKYO ENG CO LTD.
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                                                                           Kanagawa T,
Furusho K;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                      99JP-00111601.
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98US-0109732P.
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93.8%;
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LTD.
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                                                                                                  Kamagata Y,
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Pred. No. 5.
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                                                                                                     Kurata
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                                                                                                     Yamada K,
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                                                                                                     Yokomaku
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Determining the concentration of a target nucleic acid, useful e.g. for detecting genetic mutations, comprises using a fluorescently labeled probe in which emission is reduced by binding to the target nucleic acid.
                                                                                                                                                                               Example 6; Page
                                                                                                                                                                              23; 55pp; English
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The invention relates to the determination of the concentration of a CC nucleic acid target, using a fluorescently labeled probe which produces CC reduced fluorescence emission when hybridised to the target nucleic acid. CC The method comprises measuring the reduction in emission caused by hybridisation. The new method is particularly used to quantify target CC nucleic acids by a real-time polymerase chain reaction, e.g. for CC quantifying microbial cells in co-cultures or symbiotic systems, for CC quantifying microbial cells in co-cultures or symbiotic systems, for CC curves of target nucleic acids to determine a Tm value. Methods of the CC curves of target nucleic acids to be quantified quickly, easily and accurately. Particularly there is no need to remove unbound probe, and no CC materials are introduced that inhibit amplification by Taq polymerase (so CC conventional PCR conditions can be used). The specificity of PCR is kept CC injeh (amplification of primer dimers is delayed), and the limit of quantitation is reduced. Complex probes are not needed, and amplification CC can be monitored in real time. The working graph for data analysis (automatically generated by a computer) has a higher correlation (CC coefficient than conventional graphs so more accurate quantitation is coefficient that conventional graphs so more accurate quantitation is the number of G(S) in each target nucleic acid notes. its corresponding invention nucleic acid probe

20 BP; 4 A; 2 C; 0 G; 14 T; 0 U; 0 Other;

8 Query Match Best Local & Matches 1345 TATATTTTATTTCC 1360 15; Conservative 93.8%; 1.0%; 0 Score 14.4; Pred. No. 5 Mismatches .3e+02; DB 1; Length 20; Indels 0 Gaps 0

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TATATTTTTTTTTCC 20

AAH43116 standard; DNA; 20 ВP

AAH43116;

19-SEP-2001 (first entry

Antisense oligo, target HDAC-2 121-141.

Antisense; histone deacetylase; HDAC-1; cell proliferation; cancer; restenosis; fungal infections; ss. HDAC-2; HDAC-4; inhibitor; psoriasis; protozoal infec infection,

Synthetic

WO200138322-A1.

31-MAY-2001.

22-NOV-2000; 2000WO-IB001881

23-NOV-1999; 99US-0167035P

(METH-) METHYLGENE INC.

Ġ Ruel R, Lavoie R, Thibault C, Abou-Khalil

M

WPI; 2001-432601/46

ARESULT 1006
AAH43116
ID AAH4311
XX AAH4311
XX AAH4311
XX IP-SEPXX Antisen
XX Antisen
XX Antisen
XX Cell pr
XX fungal
XX Fungal
XX Synthet
XX W020013
XX W020013
XX W020013
XX IP-N
XX IP-N New inhibitors of histone deacetylase e.g. N-hydroxy-5-(4-(benzenesulfonylamino)-phenyl)-4-yn-2-pentanamide for trea for treating cancer,

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RESULT 1007
AAC89545
ID AAC8954
XX AAC8954
XX AAC8954
XX DB Human H
XX Histone
KW Histone
KW Homo 88
XX WO20007
XX WATH-:
XX MATH-:
XX MATH-:
XX Antise:
PT Macleot
PT Macleot
PT Macleot
PT Macleot
PT Deacet
PT Antise:
PT
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Best Local Similarity
Marches 15; Conserv
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Best Local Similarity
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                                                                                                                                                                                           The present invention provides inhibitors of histone deacetylase enzymes such as HDAC-1, HDAC-2, HDAC-3, HDAC-4, HDAC-5, HDAC-C and HDAC-D. These inhibitors may be antisense strands or they may be compounds identified by contacting the enzyme with the compound and measuring the resulting enzyme activity. These inhibitors are useful for treating cancers and for
                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense oligonucleotide that inhibits expression of a histone deacetylase, useful for treating and/or alleviating the symptoms of neoplasia, or for inhibiting neoplastic cell growth in an animal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-016407/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Macleod AR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Histone deacetylase; HDAC-1; HDAC-2; HDAC-3; HDAC-4; HDAC-5; HDAC-C; HDAC-C; cell cycle; tumourigenesis; cancer; inhibitor; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human HDAC-2 antisense sequence SEQ ID NO: 15.
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                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 24; 125pp; English.
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                                                                                                                                                                dentifying which histone deacetylase is involved in a neoplasia
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Conservative
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                                                                                                               A; 7 C; 7 G; 4 T; 2 U; 0 Other;
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                          1.0%;
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                          Score 14.4; DB 1;
Pred. No. 5.3e+02;
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Pred. No. 5.3e+02;
  Mismatches
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                                                    Length 20;
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Homo sapiens

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RESULT 1008
RESULT 1008
RAC89536
ID AAC8953
AC AAC8953
AC AAC8953
AC AAC8953
AC Human H
XX Histone
KW Histone
KW HAC-D;
KW Gene th
XX
Homo sa
XX
O3-MAY-
PM 03-MAY-
PR 03-MAY-
P
AAH91976/c
ID AAH9197
XX AAH9197
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention provides inhibitors of histone deacetylase enzymes such as HDAC-1, HDAC-2, HDAC-3, HDAC-4, HDAC-5, HDAC-C and HDAC-D. These inhibitors may be antisense strands or they may be compounds identified by contacting the enzyme with the compound and measuring the resulting enzyme activity. These inhibitors are useful for treating cancers and for identifying which histone deacetylase is involved in a neoplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Antisense oligonucleotide that inhibits expression of a histone deacetylase, useful for treating and/or alleviating the symptoms neoplasia, or for inhibiting neoplastic cell growth in an animal
                                                   Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis; single nucleotide polymorphism; SNP; chromosome 19p13; paternity test; chromosome 5q31-33; forensic test; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene therapy; PCR
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                                                                                                                                                                             Human inflammatory bowel disease associated polymorphic site #1051.
                                                                                                                                                                                                                                       09-OCT-2001
                                                                                                                                                                                                                                                                                                 дан91976;
                                                                                                                                                                                                                                                                                                                                                           AAH91976 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a method for detecting the presence of polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to detect the presence of genetic polymorphisms associated with inflammatory bowel disease and correlating their occurrence with disease states. They may be used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                               Recognition system; screening; identification; pharmaceutical; toxin; plant protection agent; toxin; venom; carcinogen; venom; teratogen; herbicide; fungicide; pesticide; beta-actin; human; ss.
                                                                                                  25-MAY-1999;
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10-APR-2000; 2000US-0196046P.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                              reverse phase triplex forming oligonucleotide; RP-TFO; protected nucleic acid sequence; PNAS; single nucleotide polymorphism; SNP; short tandem repeat; cancer; Factor V Leiden SNP; ss.
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                                                                                                       aminopurine
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Pred. No. 5.3e+02;
                                                                                                                                                                                                                                                                                            aminopurine
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Matches
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                                                                                          The sequence is a second reverse phase triplex forming oligonucleotide, RP-TFO (3' to the SNP) used to analyse Factor V Leiden SNP using the method of the invention. The invention relates to analysing target nucleic acid sequences comprising restricting isolated DNA, hybridising at least one triplex forming oligonucleotide (TFO), adding a 3' to 5' exonuclease to form a protected nucleic acid sequence (PNAS) tail structure, hybridising the captured structure with a single nucleotide polymorphisms (SNP) identification probe and determining the SNP score. The methods can be used for analysing target nucleic acid sequences, specially genomic DNA sequences, to determine if they contain SNPs or use in population genetics, drug development, forensics, cancer, genetic disease research, genetic analysis, diagnostics and therapeutics in
                                                                                                                                                                                                                                                                                                                                                                                          03-NOV-1999;
21-DEC-1999;
07-JUL-2000;
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                                                                                                                                                                                                                                                                     Analyzing target nucleic acid sequences, useful for population genetics, drug development and diagnosing cancer, comprises hybridizing triple forming oligonucleotide and probe to target sequence.
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                                          Sequence
                                                               Revised
                                                                                                                                                                                                                                                 Example 2; Page 66; 141pp; English.
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  15; Conserv
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OSTE C
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99US-0163416P.
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RESULT 1013
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AC
ACF57337;
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DT 16-OCT-2003 (first entry)
XX
DT Human atlastin exon 10 intron
XX
KW Human; atlastin; chromosome 1
KW HSP; neuroprotective; gene th
XX
OS Homo sapiens.
OS Synthetic.
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24-AUG-1999;
30-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention relates to a method for measuring nucleic acids using a nucleic acid probe labelled with a fluorochrome. The nucleic acid probe decreases the fluorescence of the fluorochrome when hybridised with a target nucleic acid, the decrease in the fluorescence is measured. The method can be used for measuring a target nucleic acid
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(KANK-)
(KEIZ-)
                                                                                                                                                   Human atlastin exon 10 intronic acceptor splice site.
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99JP-00242693
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                                                                             chromosome 14; 14q22.1; hereditary tive; gene therapy; intronic splice
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Pred. No. 5.
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                                                                             spastic paraplegia;
site; gene; ds.
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RESULT 1014
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KW antiinf
KW antiast
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes human atlastin, which is located to chromosome 14 (more specifically to 14922.1). Also described: (1) an isolated atlastin polypeptide; (2) identifying subjects who have, or are at risk of developing, hereditary spastic paraplegia (HSP); (3) a kit for determining if a subject has, or at risk of developing, HSP; (4) a computer readable medium encoding a representation of the atlastin nucleic acid sequence or polypeptide; (6) identifying subjects at risk of carrying an allele for HSP, and (7) treating a patient with HSP. Atlastin has neuroprotective activity and can be used in gene therapy. The atlastin nucleic acid is useful for preparing a composition for treating HSP or for identifying subjects who have, or at risk of developing, HSP. The present sequence represents an atlastin intronic splice site oligonucleotide, which is given in an example from the present invention
                                                                                                                                                                                                                                     Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New atlastin gene, useful for preparing a composition for treating Hereditary Spastic Paraplegia (HSP) or for identifying subjects who have, or at risk of developing, HSP.
 24-APR-2001; 2001US-0286137P
                                     23-APR-2002; 2002WO-US013135.
                                                                           31-OCT-2002.
                                                                                                             WO200285308-A2.
                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                   antisense gene therapy; respiratory; lung; adenosine sensitivity;
adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                              Human oligonucleotide sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                        ABZ86068 standard; DNA;
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12-SEP-2002; 2002US-00242008.
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                                                                                                                                                                                  respiratory disease; ds
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Pred. No. 5.3e+02
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                                                                                                                                                                                                  Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene thera antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allerg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               lung inflammation, lung allergies, or a respiratory disease or condition.

Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published not sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
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                                                                                                                                                    Homo sapiens
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23-APR-2002; 2002WO-US013135.

WO200285308-A2

24-APR-2001; 2001US-0286137P

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RESULT 1016
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ID ABZ8887
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Best Local
                                                                                                                                                                                                           Human; antiBenBe; lung dyBfunction; naBal Birway dyBfunction; antiInflammatory Bteroid; ubiquinone; antiInflammatory; antiBellergic; antiBellergic; antiBellergic; immunoBuppreBBive; cytostatic; gene therapy; antiBenBe gene therapy; respiratory; lung; adenoBine BenBitivity; antiBenBe gene thorapy; respiratory; lung; adenoBine BenBitivity; adenoBine receptor; bronchodilation; bronchoconBtriction; lung allergy;
                                                                                                                                                                                                                                                                                                                               Human oligonucleotide sequence.
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 24-APR-2001; 2001US-0286137P
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                                         23-APR-2002; 2002WO-US013135.
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The invention relates to a novel pharmaceutical composition, which has a Cfirst active agent comprising an oligonucleotide antisense to the Cinitiation codon, coding region, 5 or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine creceptor, producing bronchodilation, increasing levels of bulquinone or lung sinflammation, lung allorgies, or a respiratory disease or condition. CC Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO CC at fry wino int/num/hilainda or emmerces.
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Sequence
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  BP;
16 A; 1 C; 1 G; 2 T; 0 U; 0 Other;
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Pred. No. 5.
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DB 1;

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Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antianflammatory steroid; ubiquinone; antinflammatory; ene therapy; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;

Human CCR3 oligonucleotide sequence.

24-APR-2001; 2001US-0286137P

23-APR-2002; 2002WO-US013135.

WO200285308-A2 Homo sapiens.

inflammation;

respiratory

disease;

dв

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RESULT 1018
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XX ABZ9151
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   24-APR-2001; 2001US-0286137P
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adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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Pred. No. 5.3e+02
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The invention relates to a novel pharmaceutical composition, which has a cfirst active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5 or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition of treating or CC preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an infiniflammatory steroid in a subject, for reducing or depleting levels of adenosine CC receptor, producing bronchodilation, increasing levels of adenosine column inflammation, lung allergies, or a respiratory disease or condition. CC Note: The sequence data for this patent is not represented in the printed capacitication, but was obtained in electronic format directly from WIPO as ferre with a first patent is not represented in the printed capacitication, but was obtained in electronic format directly from WIPO
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  Matches
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                                                                                          Sequence
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  15;
                       Similarity
                                                                                          20 BP; 2 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
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  Conservative
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                       93.8%;
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Pred. No. 5
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    Mismatches
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PR 23-J-PR
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                                                                                                                                                                                           Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; pypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
24-APR-2001; 2001US-0286137P
                                   23-APR-2002; 2002WO-US013135
                                                                                                                                                                                                                                                                                                        Human oligonucleotide sequence.
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                                                                        31-OCT-2002
                                                                                                         WO200285308-A2
                                                                                                                                            Homo sapiens
                                                                                                                                                                               inflammation;
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RESULT 1020
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KW antiinf
KW antiest
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Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
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Best Local S
Matches 15
                                                                                                                                                                                                   Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; antisense gene thorapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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   24-APR-2001; 2001US-0286137P
                                                                                                                                                                                                                                                                                                                     Human oligonucleotide sequence
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                                      23-APR-2002; 2002WO-US013135.
                                                                                                                WO200285308-A2
                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of
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                                                                                                                                                                                     inflammation;
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, Shahabuddin
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Pred. No. 5
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Sequence 20

BP; 5 A; 2 C; 2

G; 11 T; 0 U; 0 Other;

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The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or CC massal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC has antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC immunosuppressive, and cytostatic activity. The composition may have a CC for enhancing the prophylactic or therapautic respiratory effect of an CC enhancing the prophylactic or therapautic respiratory effect of an CC or reducing sensitivity to adenosine, reducing or depleting levels of adenosine creceptor, producing bronchodilation, increasing levels of adenosine creceptor, producing bronchodilation, increasing levels of bronchoconstriction, cC lung inflammation, lung allergies, or a respiratory disease or condition. CC Note: The sequence data for this patent is not represented in the printed confective to the composition of the c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO 8778; 872pp; English
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                                                         ftp.wipo.int/pub/published_pct_sequences
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L, Shahabuddin
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             1295
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        TGGTTAATCTATTTTT 1310
TGGTTAATCTTTTTTT
                          Conservative
                                 1.0%;
16
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                                 Score 14.4;
Pred. No. 5.
                           Mismatches
                                .3e+02;
                                         DB
                                        Length
                           Indels
                                         20
                           0,
                           Gaps
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RESULT 1021
ABZ88813
ID ABZ8881
XX ABZ8881
XX ABZ8881
XX IT ABZ8881
XX ABZ881
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Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antialsthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; inflammation; respiratory

sapiens

WO200285308-A2

23-APR-2002; 2002WO-US013135.

24-APR-2001; 2001US-0286137P

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RESULT 1022
ABZ86072
ID ABZ8607
XX ABZ8607
XX ABZ8607
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XX Human,
KW antiinf
KW antiast
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antisfialmentory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO as for whis patent is not represented in the printed as pecification, but was obtained in electronic format directly from WIPO as for many patent is not represented in the printed as pecification, but was obtained in electronic format directly from WIPO as for many patent is not represented in the printed as pecification, but was obtained in electronic format directly from WIPO as format directly from WIPO as the many patent is not represented in the printed as pecification.
                                                                                                                                                                                                                                                        antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                 23-APR-2002; 2002WO-US013135
                                                                                                                                                                                                                                                                                                                                                                                               Human oligonucleotide sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; SEQ ID NO 4055; 872pp; English
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    24-APR-2001; 2001US-0286137P
                                                                                                31-OCT-2002.
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                                                                                                                                                                                                                                     inflammation;
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L, Shahabuddin
                                                                                                                                                                                                                                     respiratory disease;
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93.8%;
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Pred. No. 5.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention
                                                                                                                                                                                                                              Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiaflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                         beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                     pulmonary transplantation rejection; ss; primer
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Pred. No. 5.3e+02
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23-APR-2002; 2002WO-US013143.

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RESULT 1024
ABD25043
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Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibros
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allers reducing adenosine sensitivity, levels of adenosine (A) or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                            AI128305-derived
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                                                                                                                                         oligonucleotide SEQ
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Pred. No. 5.3e+02;
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                                                            inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary distression, emphysema, chronic obstructive pulmonary cransplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to
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       This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating composition, respiratory tract inflammation, allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, composition of allergies are derived from a gene encoding or regulating complete tides are derived from a gene encoding or regulating composition or cancer and can be anti-sense to the corresponding mRNA. Composition also describes a kit, that comprises: (a) a delivery composition also describes a kit, that comprises: (a) a delivery composition for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, composition has antiallergic, antiinflammatory, antiasthmatic, composition is useful for preventing or creating a respiratory, lung or malignant disease. The administered composition composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung and composition and/or lung.
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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptc surfactant depletion or hyposecretion, when administered to a mammal. oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA.
                                                                                                                                                                                                               Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; enalgesic; hypotensive; dimmunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypotension; emphysems; chronic obstructive pulmonary disease; cancer; bronchitis;
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                                                                                                                                                            Claim 15; SEQ ID NO 12949; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                               24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-APR-2002; 2002WO-US013143.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pulmonary transplantation rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human CCR3-derived oligonucleotide SEQ ID 12949.
                                                                                                                                                                                               bronchodilating agent.
                                                                                                                                                                                                                                                                                       WPI; 2003-093058/08
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                                                                                                                                                                                                                                                                                                                                                                              (EPIG-) EPIGENESIS PHARM INC
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ilarity 93.8%;
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, Shahabuddin
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S;
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RESULT 1028
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CC distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system ce.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
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Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targunucleic acids associated with lung airway or lung dysfunction
                                                                                                                                                                                                                                    23-APR-2002; 2002WO-US013143.
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                                                                                                              Sandrasagra A,
L, Shahabuddin
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Pred. No. 5.3e+02;
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targeted to

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CC comprising oligonucleotides, effective for alleviating
CC bronchoconstriction, respiratory tract inflammation, allergies and
CC reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors,
CC surfactant depletion or hyposecretion, when administered to a mammal. The
CC eligonucleotides are derived from a gene encoding or regulating
CC expression of a target polypeptide associated with lung airway or lung
CC dysfunction or cancer and can be anti-sense to the corresponding mRNA.
CC The invention also describes a kit, that comprises: (a) a delivery
CC device, in separate containers, (b) the oligonucleotides, (c)
CC instructions for adding a carrier and for use of the kit. The composition
CC of the invention has antiallergic, antiinflammatory, antiasthmatic,
CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a
CC treating a respiratory, lung or malignant disease. The administered
CC composition comprises oligo and is administered to reduce the production
CC or availability, or to increase the degradation of the target mRNA or to
CC reduce the amount of target polypeptide present in the lungs. The
CC inflammation, allergies and/or surfactant hypoproduction are associated
CC with a disease or condition such as pulmonary vasoconstriction,
CC distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary
CC distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary
CC distress syndrome, pain, cystic fibrosis, allergic rhinitis or cancer.
CC the reduced adenosine content of the anti-sense oligos corresponding to
CC thymidines present in the target RNA serves to prevent the breakdown of
CC the oligonucleotides into products that free adenosine into the system
CC the oligonucleotides into products that free adenosine into the system
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CC the oligonucleotides into green the system
CC the oligonucleotides into field the content of the coligonucleotides content to it
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                                                                                                                                                                                                        respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                        23-APR-2002; 2002WO-US013143
                                                                                                                                                                                                                                                                                                                                                                                                                     Human stanniocalcin-derived oligo SEQ ID 1314.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABD22302 standard;
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Pred. No. 5.3e+02;
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RESULT 1030
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CC pronchoconstriction, respiratory tract inflammation, allergies and
CC reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors,
CC surfactant depletion or hyposecretion, when administered to a mammal. The
CC oligonucleotides are derived from a gene encoding or regulating
CC expression of a target polypeptide associated with lung airway or lung
CC dysfunction or cancer and can be anti-sense to the corresponding mRNA.
CC The invention also describes a kit, that comprises: (a) a delivery
CC device, in separate containers, (b) the oligonucleotides, (c)
CC instructions for adding a carrier and for use of the kit. The composition
CC of the invention has antiallergic, antiinflammatory, antiasthmatic,
CC beta-adrenergic agonist. The composition is useful for preventing or
CC treating a respiratory, lung or malignant disease. The administered
CC composition comprises oligo and is administered to reduce the production
CC or availability, or to increase the degradation of the target mRNA or to
CC reduce the amount of target polypeptide present in the lungs. The
CC inflammation, allergies and/or surfactant hypoproduction are associated
CC with a disease or condition such as pulmonary vasconnstriction.
CC inflammation, allergies, asthma, impeded respiration, respiratory
CC thymidines present in the target RNA serves to pronchitis or cancer.
CC thymidines present in the target RNA serves to prevent the breakdown of
CC thymidines present in the target RNA serves to prevent the breakdown of
CC the oligonucleotides into products that free adenosine into the system
CC entered adenosine content of the anti-sense oligos corresponding to
CC thymidines present in the target RNA serves to prevent the breakdown of
CC the oligonucleotides into products that free adenosine into the system
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Miller S
             Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibros
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                        AA102454-derived
                                                                                                                                                                 29-JUL-2004
                                                                                                                                                                                                           ABD27748;
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L, Shahabuddin
                                                                                                                     oligonucleotide SEQ ID 6760.
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                   fibrosis;
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RESULT 1031
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Best Local
                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  comprising oligonuclectides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) recepts surfactant depletion or hyposecretion, when administered to a mammal oligonuclectides are derived from a gene encoding or regulating
                                                                                                                                                                                                                                                                                                                        inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 15; SEQ ID NO 6760; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-APR-2001; 2001US-0286036P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a novel composition (a) a first active agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-APR-2002; 2002WO-US013143.
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                                                                                                                                                                                                   Local
                                                                                                                         666 GACTCACCTCTGACGC 681
                                                                                                                                                                                    15; Conserv
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                                                                                                                                                                                                                                                                   20
                                                                                                   GACTCACCTCTGTCGC 16
                                                                                                                                                                                                                                                                BP; 2 A; 7 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                         unwanted effects due
                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sandrasagra A, K
L, Shahabuddin S;
                                                                                                                                                                                                   1.0%;
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                                                                                                                                                                                                       Score 14.4;
Pred. No. 5.
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                                                                                                                                                                                                   5.3e+02;
                                                                                                                                                                                                                        DB 1;
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                                                                                                                                                                                                                      Length 20
                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (A) receptors,
                                                                                                                                                                                    0;
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RESULT 1032
ADK80880/c
ID ADK8088
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AC ADK8088
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DT 20-MAY-
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                                                                                                                                                                                                                                                                                           The present invention relates to an oligonucleotide anti-sense to e.g., CC initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end funcleot acid target comprising gene(s) chosen from e.g. CC end of nucleic acid target comprising gene(s) chosen from e.g. CC oligonucleotide and optionally surfactant operatively linked to the coligonucleotide. The method is useful for preventing or treating a cC oligonucleotide. The method is useful for preventing to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is co f a subject an effective amount of an inhibitor. The oligonucleotide is co f a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded consent for constructive pulmonary diseases conservation, cystic fibrosis (F), chronic obstructive pulmonary diseases conservation, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the conservation.
                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 2; SEQ ID NO 420; 85pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-JUL-2002; 2002US-0399076P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligonucleotide associated to CCR3 #65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADJ59564;
                                                                                                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-JUL-2003; 2003WO-US023509.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                            1524 AAAAAAAAGTAAAAGG 1539
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                                                                                                                                                                17
                                                                                                                                                                                                                            15;
                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                AAAAAAAGTACAAGG
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lin S,
                                                                                                                                                                                                                                                                                         B₽;
                                                                                                                                                                                                                            Conservative
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Lu H, Cong H;
                                                                                                                                                                                                                                                                                         2 A; 4 C; 2 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                           1.0%;
93.8%;
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Pred. No. 5
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ADK80880;

ADK80880 standard;

DNA;

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ВÞ

20-MAY-2004

(first entry)

Chimeric

phosphorothioate oligonucleotide to target Nav1.3 #8214.

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RESULT 1033
AD045054/c
ID AD04505
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AC AD04505
XX
AC AD04505
XX
DT 15-JUL-
XX
DT 15-JUL-
CKW Human c
XX
CCRL; c
KW CCRL; c
KW CCRL; c
KW Ltyptas
KW Lung di
KW asthma;
KW asthma;
KW asthma;
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Best Local S
Matches 15
Human; 58; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsivenese; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; COPD; allergic rhinitis; accute respiratory distress syndrome; pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target different regions of the human Navl.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     wew antisense compound targeted to a nucleic acid molecule encoding Navl.3, useful for useful for treating a disease or condition associated with Navl.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acid molecule encoding Nav1.3, where the antisense compound specifically hybridizes with and inhibits the expression of Nav1.3. The compound and composition are useful for treating a disease or condition
                                                                                                                                                                      Human oligonucleotide #420
                                                                                                                                                                                                            15-JUL-2004
                                                                                                                                                                                                                                                 ADO45054;
                                                                                                                                                                                                                                                                                   ADO45054 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PHAA
                                                                                                                                                                                                                                                                                                                                                                                                    1252 TTTTGTTTTTTAATCA 1267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2004-203785/19.
                                                                                                                                                                                                                                                                                                                                                                           17 TTTTGATTTTTAATCA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; SEQ ID NO 8214; 417pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    l Similarity
15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention relates to an antisense compound
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; 2 C; 3 G; 4 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1.0%;
93.8%;
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Pred. No. 5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
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RESULT 1034 ADO55869/c ID ADO5586

ADO55869 standard; DNA;

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                                                                                    Query Match
Best Local
                                                                      Matches
                                                                                                                                                                                          allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction obsonchoconstriction. This sequence represents an oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel single or multiple target oligonucleotide anti-sense to e.g. Coninitiation codon, intron of respiratory disease-relevant gene e.g. CORANTES, MCP4, useful for prophylaxis or treating respiratory disease
                                                                                                                                        Sequence 20 BP; 2 A; 4 C; 2 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-APR-2002; 2002WO-US013135.
23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 2; SEQ ID NO 420; 174pp; English.
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Shahabuddin S, Lu H, Cong H;
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                                                                                                                                                                             invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lung inflammation; bronchitis; airway obstruction; bronchoconstriction
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SHAH/)
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                                 1524 AAAAAAAAAGTAAAAGG 1539
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                                                                                      Similarity
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SHAHABUDDIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGUILAR D.
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 AAAAAAAGTACAAGG
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                                                                    Conservative
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                                                                                    1.0%;
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                                                                                    Score 14.4;
Pred. No. 5
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                                                                                    .3e+02
                                                                                                       DB 1; Length
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ADO55807
ID ADO5580
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AC ADO5580
XC ADO5580
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC The present invention relates to antisense compounds targeted to a control of control of the control of con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  never in mitosis gene a-related kinase 6; hyperproliferative disorder; cancer; cytostatic; ds.
Antisense therapy; human; NIMA-related never in mitosis gene a-related kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human NIMA-related kinase 6 DNA target sequence
                                                                                                                     Human NIMA-related kinase 6 DNA, antisense oligonucleotide
                                                                                                                                                                                                                                                                              AD055807;
                                                                                                                                                                                                                                                                                                                                                           ADOSS807 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-NOV-2002; 2002US-00295471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US2004097441-A1
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                                                                                                                                                                                                  12-AUG-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2004-389184/36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AGTGGGCGGCGGTGGT 1051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 115;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2002US-00295471.
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                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; 9 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            51pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 14.4; D
Pred. No. 5.3e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                English.
    | kinase 6;
| 6; hyperp:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5.3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
    hyperproliferative disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20;
                                                                                                                         #30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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RESULT 1036
ADP20520/c
ID ADP2052
XX
AC ADP2052
XY
AC ADP2052
XX
DT 26-AUG-
XX
Transcr
XX
KW cytosta
KW AP-2 al
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                                                                                                                                                                                                                                Ś
                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to antisense compounds targeted to a nucleic acid encoding human never in mitosis gene a related kinase 6 (NIMA-related kinase 6). The antisense compound comprises an antisense oligonucleotide that specifically hybridises with the nucleic acid and inhibits the expression of NIMA-related kinase 6. The antisense oligonucleotide is a chimeric oligonucleotide. The antisense oligonucleotide comprises at least one modified internucleoside linkage, preferably a phosphorothioate linkage. It also comprises at least one modified sugar molety, preferably a "O-methoxyethyl (2'-MOE) sugar molety. The antisense oligonucleotide further comprises at least one modified nucleobase, preferably a 5-methylycosine. The antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides for modulating never in mitosis, gene (NIMA)-related kinase 6 expression, useful for diagnosing, preventing treating diseases associated with the kinase, e.g. hyperproliferative
                                                                                                                                                                                                                                                                                                                             Sequence 20
                                                                                                                                                                                                                                                                                                                                                          present invention.
                                                                                                                                                                                                                                                                                                                                                                        oligonucleotides are useful for the treatment of diseases such as hyperproliferative disorders, e.g. cancer. The present sequence represents an antisense oligonucleotide used in the examples of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dobie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-NOV-2002; 2002US-00295471.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer; cytostatic; phosphorothioate;
                                            Transcription
                                                                           26-AUG-2004
                                                                                                       ADP20520
                                                                                                                                  ADP20520 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-MAY-2004.
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                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                        1036
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                                                                                                                                                                                                           \vdash
                                                                                                                                                                                                                                                                    l Similarity 93.8
15; Conservative
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                                                                                                                                                                                                                                     AGTGGGCGGCGGTGGT 1051
                                                                                                                                                                                                           AGTGGGCTGCGGTGGT 16
                                                                                                                                                                                                                                                                                                                              B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 44; 51pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002US-00295471.
                                            factor AP-2 antisense oligonculeotide seqid 67.
                                                                           (first
                                                                                                                                                                                                                                                                                                                              2 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note "This oligonucleotide has a phosphorothioate backbone and 2'-methyoxyethyl (2'-MOE) wings at the and 3' ends, which are 5 nucleotides in length at ea end. All cytidine residues are 5-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
1. .20
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                                                                         entry)
                                                                                                                                                                                                                                                                                1.0%;
                                                                                                                                                                                                                                                                                                                              4
C;
                                                                                                                                                                                                                                                                                                                              9 G;
                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                  Score 14.4; DB 1
Pred. No. 5.3e+02
0; Mismatches
                                                                                                                                                                                                                                                                                                                              5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  88.
                                                                                                                                                                                                                                                                                                                                0 Other
                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                     Indels
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cytostatic; AP-2-Inhibitor-Alpha; AP-2 alpha; AP-2 alpha modulator; AP-2 alpha associated disorder; hyperproliferative disorder; human;

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ADP21858/c
ID ADP2185
XX ADP2185
AC ADP2185
XX ADP2185
XX ADP2185
XX 26-AUG-
XX Cytosta
XW cytosta
XW cytosta
XW ornithi
XW hyperpr
XX
OS Homo sa
XX
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modulator of AP-2 alpha by contacting a preferred target segment of a nucleic acid molecule encoding AP-2 alpha with one or more modulatere modulators of AP-2 alpha, and identifying one or more modulators of AP-2 alpha with one candidate modulators of AP-2 alpha, and identifying one or more modulators of AP-2 alpha expression, which modulate the expression of AP-2 alpha; a diagnostic method for identifying a disease state; and a kit or assay device comprising (I). The compound is useful for treating an animal having a disease or condition associated with AP-2 alpha, particularly a hyperproliferative disorder. The compounds may be used for diagnostics, therapeutics prophylaxis and as research reagents; or as tools in differential and/or combinatorial analyses to elucidate expression patterns of a portion or the entire complement of genes expressed within cells and tissues. This sequence represents a human transcription factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention describes a compound (I) 8-80 nucleobases in length targeted to a nucleic acid molecule encoding AP-2 alpha. The compound specifically hybridises with a nucleic acid molecule encoding AP-2 alpha (19868 bg, SEQ ID NO: 4), and inhibits the expression of AP-2 alpha. Also described are:inhibiting the expression of AP-2 alpha in cells or tissues comprising contacting the cells or tissues with (I); screening for a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New compounds targeted to nucleic acid molecules encoding AP-2 alpinibits the expression of AP-2 alpha, useful for treating AP-2 alassociated disease or condition, particularly a hyperproliferative
                    Homo sapiens
                                                      hyperproliferative
                                                                         cytostatic; gene therapy; ornithine decarboxylase 1; ornithine decarboxylase 1 associated disorder;
                                                                                                                              Human ornithine
                                                                                                                                                                    26-AUG-2004
                                                                                                                                                                                                      ADP21858
                                                                                                                                                                                                                                          ADP21858 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 15; SEQ ID NO 67; 58pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-DEC-2002; 2002US-00315962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-DEC-2002; 2002US-00315962
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2004109848-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AP-2 antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISIS-) ISIS PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 5 A; 6 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             factor; antisense oligonucleotide; antisense technology;
                                                                                                                              decarboxylase 1
                                                        disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                        1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Freier SM,
                                                                                                                                                                                                                                            ΒP
                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                        cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.4; DB 1;
Pred. No. 5.3e+02;
                                                                                                                              primer seqid 6.
                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                      human; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                             20
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention describes a new compound, having a sequence comprising 8-80 bp targeted to a nucleic acid encoding ornithine decarboxylase 1, specifically hybridises with the nucleic acid encoding ornithine decarboxylase 1 comprising 2035-bp sequence and inhibits expression of ornithine decarboxylase 1. Also described are: inhibiting the expression of ornithine decarboxylase 1 in cells or tissues; screening for a modulator of ornithine decarboxylase 1 identifying a disease state; a kit or assay device comprising the compound; and treating an animal having a disease or condition associated with ornithine decarboxylase 1. The oligonucleotide compound is useful for preparing a composition for treating hyperproliferative disorder, e.g. cancer. This sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New oligonucleotide compound that inhibits expression of ornithine decarboxylase 1, useful for preparing a composition for treating hyperproliferative disorder, e.g. cancer.
                           WPI; 1995-018287/03.
                                                                              16-APR-1993;
                                                                                                       16-APR-1993;
                                                                                                                                 01-NOV-1994
                                                                                                                                                          JP06303997-A
                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                           aggregate;
                                                                                                                                                                                                                          Analysis;
                                                                                                                                                                                                                                                Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                            04-AUG-1995
                                                                                                                                                                                                                                                                                                     AAQ75552;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              decarboxylase 1.
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                                                    (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                   729 TGTTGCTGCTGCCTTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                refrectecter 5
                                                                                                                                                                                                             gene expression; reverse transcription; primer; cDNA;
; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                              93JP-00112515
                                                                                                        93JP-00112515
                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.0%;
                                                                                                                                                                                                                                                                            entry)
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Pred. No. 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB
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Analysis of cDNA and gene expression -

by amplification of

mRNA followed

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RESULT 1039
ADL79331
XX ADL7933
X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                            20-FEB-2002;
11-MAR-2002;
29-MAY-2002;
06-JUN-2002;
06-JUN-2002;
03-JUL-2002;
29-AUG-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A method for the analysis of cDNA comprises (a) preparing an aggregate (double-stranded cDNAs by using an aggregate of mRNAs and a plural type (labelled reverse transcription primers (GENESEQ files AAO75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; EGFR; HER2; EGFR2; neu; erbB2; c-erb-B-2; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA interference; short interfering nucleic acid; siNA; short interfering RNA; double-stranded RNA; micro-RNA; miRNA; short interfering RNA; skRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification;
                                                          New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the epidermal growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human HER2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 5; llpp; Japanese.
                                                                                                                                                                                                                                                         (RIBO-)
                                                                                                                                                                                                                                                                                                                                                                       19-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1246 TCTTTGTTTTGTTTTTAA 1264
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                               receptor gene
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                                                                                                                                                                                                                                                                                                          2002US-0393924P.
2002US-0406784P.
2002US-04083789.
2002US-0409293P.
2002US-00251117.
2002US-00277494.
2003US-0440129P.
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2002WO-US016840.
2002US-00163552.
2002US-0386782P.
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                                                                                                                                                                                                                                                         INC.
                                                                                                                                                                                                  Beigelman
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Pred. No. 6.
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                                                                                                                                                                                                  Fosnaugh K,
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CC interference. The siNAs may or may not comprise ribonucleotides and may CC be double or single stranded. They further comprise sense and antisense CC regions, or alternatively are assembled from a sense oligonucleotide and CC an antisense oligonucleotide. Specifically, the siNAs include short CC interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short CC can contain deoxyribonucleotides, and can be unmodified or chemically modified, CC can contain deoxyribonucleotides, and can be chemically synthesised, CC expressed from a vector or enzymatically synthesised. The invention also CC relates to kits for the in vitro or in vivo delivery of siNA; conjugates CC and/or complexes of siNA; and vectors that express siNA. The siNAs are CC used to modulate expression of EGFR genes in cells, tissue explants or CC organisms (e.g., by ex vivo gene therapy), or in grafts and transplants CC for the treatment of a variety of conditions. They may be used for treating a wide range of cancers such as breast and ovarian cancer. The CC identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide CC polymorphisms). The present sequence represents the lower strand of a CC HER2 (EGFR2)-targeted double-stranded siNA.
                    Query Match
                                                                         Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          downregulate expression receptor (EGFR) genes (
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Local
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Similarity
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                                                                       BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          elates to short interfering nucleic acids (siNA) which pression of one or more human epidermal growth factor genes (including HER1, HER2 HER3 and HER4) by RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NO 496; 171pp; English.
                                                                       P
                                                                       2 C; 0 G; 0
Score 14.2; DB 1;
Pred. No. 6.1e+02;
                                                                       T; 1 U;
                                                                         0 Other;
                         Length 19;
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Query Match

1.0%; Score 14.2; DB 1; Length 19;

Best Local Similarity 78.9%; Pred. No. 6.1e+02;

Matches 15; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

Qy 1519 TAAAAAAAAAAAAAA 1537

Qb 1519 TAAAAAAAAAAAAAA 1537

Db 1 UAAAAAAAAAAAAAAA 19

ADL79082/c
ID ADL790
XX
AC ADL790
AC 06-JUN-2002 06-JUN-2002; 03-JUL-2002; 03-JUG-2002; 05-SEP-2002; 09-SEP-2002; 19-SEP-2002; 21-OCT-2002; 20-FEB-2002; 11-MAR-2002; 29-MAY-2002; RNA interference; short interfering nucleic acid; siNA; miRNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; pharmacogenomics; gene function analysis; gene mapping; cancer; cytostatic; human; oncogene; epidermal growth factor receptor; EGFR; cytostatic; 20-FEB-2003; 28-AUG-2003. WO2003070912-A2 Human HER2 (EGFR2) transcript target sequence/siNA upper strand, SEQ:247. 20-MAY-2004 ADL79082; ADL79082 standard; RNA; 19 sapiens. EGFR2; neu; erbB2; c-erb-B-2; target sequence; ss. 2002US-0408378P. 2002US-0409293P. 2002US-00251117. 2002US-00277494. 2002US-00163552. 2002US-0386782P. 2002US-0393924P. 2002US-0406784P. 2002US-0358580P. 2002US-0363124P. 2003WO-US005045 2002WO-US016840. (first entry) ВP

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RESULT 1041
ABZ89546/c
ID ABZ8954
XX ABZ8954
XX ABZ8954
XX IT-OCT-
XX Human o
XX Human o
XX Human;
KW Antiast

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                     Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
         23-APR-2002; 2002WO-US013135
                                                                      31-OCT-2002.
                                                                                                                                    WO200285308-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human oligonucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABZ89546;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABZ89546 standard; DNA;
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receptor (EGFR) genes (including HER1, HER2 HER3 and HER4) by RNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-697612/66.
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                                                                                                                                                                                                                                                              inflammation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                           respiratory disease; ds
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downregulates
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Pred. No. 6.1e+02;
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RESULT 1042
ABD25776/c
ID ABD2577
XX ABD2577
XX ABD2577
XX ABD2577
XX ABD2577
XX Human;
XW respirac
XW surfact
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine rung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                 Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; allergic rhinitis; pulmonary hyportension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 18 A; 0 C; 0 G; 2 T; 0 U; 0 Other;
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L, Shahabuddin
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WO200285309-A2 Homo sapiens pulmonary transplantation

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RESULT 1043
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AC AAQ7557
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DT 04-AUG-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mNNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or burnchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated inflammation, allergies and/or surfactant hypoproduction are associated
                                                                                                                                                                                                                                                                                                                                                                                                                                       with a disease or condition such as pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system
Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                         the oligonucleotides into products that e.g., lung, brain, heart, kidney, etc, t prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     surfactant depletion or hyposecretion, when administered to a mammal. oligonuclectides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) rec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 15; SEQ ID NO 4788; 763pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nucleic acids associated bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
                                                 Reverse transcription primer used in cDNA analysis technique.
                                                                                    04-AUG-1995
                                                                                                                    AAQ75579
                                                                                                                                                      AAQ75579 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel composition
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                                                                                                                                                                                                                                                                                                                        Local Similarity
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L, Shahabuddin
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 A method for the analysis of cDNA comprises (a) preparing an aggregate
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RESULT 1044
AAQ75582
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Best Local Similarity
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                             Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                                                               16-APR-1993;
                                                                                                                                                                                                                         JP06303997-A
                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                    Analysis; gene expression; reverse transcription; primer; cDNA, aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                            Reverse transcription primer used in cDNA analysis technique.
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by digestion with restriction enzymes.
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Disclosure; Page 5; 11pp; Japanese
                                                                          WPI; 1995-018287/03.
                                                                                                      (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                   16-APR-1993;
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                                                                                                                                                                                                                                                                                   aggregate;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               can be used to analyse gene expression rapidly and easily
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RESULT 1045
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                                                                                                                                                                                                                       Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Analysis of cDNA and gene expression - by digestion with restriction enzymes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    restriction enzyme;
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                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93JP-00112515.
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                                                                                                                                     84.2%;
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/me; ss.
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                                                                                                                                     Score 14.2; DB 1
Pred. No. 5.8e+02
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Pred. No. 5.8e+02;
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RESULT 1046 ABZ88694/c

RESULT 1047 AAQ75596

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the contribution codon, coding region, 5' or 3' end genomic flanking regions, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of contributions of genes encoding a polypeptide associated with lung and/or consal airway dysfunction and a second active agent comprising an cantinflammatory steroid and ubsquinone. A composition of the invention continuous pressive, and cytostatic activity. The composition may have a constituent of the invention continuous pressive, and cytostatic activity. The composition may have a consecution are presented as a consecution of useful for treating or preventing a respiratory, lung or malignant disease or condition, also continuous pressive, and cytostatic activity. The composition may have a consecution are prophylactic or therapeutic respiratory effect of an antisense gene therapy. The composition is useful for treating or preventing a respiratory before the invention or reducing levels of adenosine continuous producing bronchodilation, increasing levels of adenosine continuous producing bronchodilation, increasing levels of bulquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, continuous producing alergies, or a respiratory disease or condition. Onte: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at the printed and continuous continuous
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                                                                                                                                                                              Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antisense; lung dysfunction; nasal airway dysfunction; antishflammatory steroid; ubiquinone; antiinflammator; hypotensive immatters.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 3936; 872pp; English.
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                                                                                                                                                                                                                     ftp.wipo.int/pub/published_pct_sequences
                                         1246
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                                                                                                        Score 14.2; DB 1;
Pred. No. 5.8e+02;
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Best Local 9
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                                                                                                                                                                                                                                                                                        aggregate;
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                              16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               alysis of cDNA and gene expression - by digestion with restriction enzymes.
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Pred. No. 5.8e+02;
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RESULT 1049
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Best Local
                          A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ filles AQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in separate lanes. The electrophoresing the digested aggregate of cDNAs in separate
Sequence 20 BP; 1 A; 1 C; 1 G; 17 T; 0 U; 0 Other;
                                                                                                                                                              Disclosure; Page 5; 11pp; Japanese.
                                                                                                                                                                                                                                         WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                   16-APR-1993;
                                                                                                                                                                                                                                                                                                                               16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Analysis; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Reverse transcription primer used in cDNA analysis technique.
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by digestion with restriction enzymes.
                                                                                                                                                                                                            Analysis of cDNA and gene expression -
                                                                                                                                                                                                                                                                     (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                              01-NOV-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    aggregate;
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                                                                                                                                                                                            digestion
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16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     restriction enzyme;
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                                                                                                                                                                                            with
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                                                                                                                                                                                                                                                                                                   93JP-00112515
                                                                                                                                                                                                                                                                                                                                 93JP-00112515
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                                                                                                                                                                                              restriction
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84.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  reverse transcription; primer; cDNA;
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Pred. No. 5.8e+02
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Query Match

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Score 14.2;

DB 1; Length

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RESULT 1050
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                                                                           The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a cuse in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of antiinflammatory steroid in a subject, for reducing levels of adenosine receptor, producing sensitivity to adenosine, reducing levels of adenosine conceptor, producing bronchodilation, increasing levels of adenosine or condition. In a subject, a tissue, or treating bronchoconstriction, constriction, ing sallegies, or a respiratory disease or condition. The sequence data for this patent is not represented in the printed as precification, but was obtained in electronic format directly from WIPO as from wipo int/nuh/nuh/sed not sequences.
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Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiguinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 15; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-229219/22.
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                                                            ftp.wipo.int/pub/published_pct_sequences
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, Tang L,
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     1 A;
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L, Shahabuddin
     1 C; 0
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        G; 18
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     0 U;
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Query Match

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RESULT 1051
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CC comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating adenosine sensitivity, levels of adenosine (A) or (A) receptors, creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, construction or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating construction of a target polypeptide associated with lung airway or lung construction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery construction of for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung confilammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, respiratory construction, and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatrory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema, chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 15; SEQ ID NO 554; 763pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sandrasagra A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein A2-derived oligo SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      a novel composition (a)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a first active
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RESULT 1052
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PN JP06303
XX O1-NOV-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                         Query Match
Best Local
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                                                                                                                                                                                                                                                                                  A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files ANQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Analysis; gene expression; reverse transcription; primer; cDNA; aggregate; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 1 A; 1 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ75586 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 5; llpp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NITE ) NIPPON TELEGRAPH & TELEPHONE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   alysis of cDNA and gene expression digestion with restriction enzymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1302
                                                      1248 TTTGTTTTTTTTAATC 1266
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                                                                                                                1 Similarity
16; Conserv
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                                                                                                                                                                                                                              20 BP; 1 A; 1 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TTTTTTTTTTTTTTCA 20
                                                                                                                    Conservative
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                                                                                                                                            84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           84.2%;
ITTTTATC 20
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                                                                                                                Score 14.2; DB 1;
Pred. No. 5.8e+02;
0; Mismatches 3;
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Pred. No. 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                       Length 20;
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RESULT 1053

30-SEP-1994

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RESULT 1054
AAQ68872
ID AAQ6887
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AC AAQ6887
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AC AAQ6887
C SS-MAR-
DT 25-MAR-
DT 31-MAY-
XX Oligonu
XX Sportasto
XX Heparit
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                   Oligonucleotide; antisense; self paired; nuclease resistant; dermatological disorders; viral infection; cancer; atopical c psoriasis; melanoma; T cell lymphoma; herpes simplex; papillo hepatitis; HIV; human immunodeficiency virus; oncogene; colla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEO files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                         elastase; bone marrow graft; ss.
                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide (SA12/ml) used as control in antisense therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
31-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ68872;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ68872 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 5; 11pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1995-018287/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-NOV-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Analysis; gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-AUG-1995
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                                                                 FR2703053-A1
                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 2 A; 0 C; 1 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
(first entry)
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Pred. No. 5.
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                                                                                                                                                                                                                                                                                       dermatitis;
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RESULT 1055
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New hooked or semi-hooked oligonucleotides (see AAQ68869-71, AAQ68873, AAQ68875, AAQ68877, AAQ68879 and AAQ68880) are useful as therapeutic antisense molecules for treating dermatological disorders (e.g. atopical dermatitis, psoriasis, melanoma, T cell lymphoma etc.) viral infections (e.g. herpes simplex, papilloma, hepatitis or HIV); or cancer (when directed against an oncogene), due to their ability to hybridise with target nucleic acid. They can be used ex vivo, e.g., to treat bone marrow grafts. They can also be used for diagnosis or in cosmetics e.g. to block mRNA coding proteins involved in the ageing process such as collagenase or elastase. This linear antisense oligonucleotide is used as a control to see whether the hooked and semi-hooked oligonucleotides exhibit a greater resistance to exonucleases than linear oligonucleotides. (Updated on 25-MAR-2003 to correct PN field.) (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New oligo:nucleotide(s) self paired at one or both ends - have improved resistance to nuclease(s) and reduced toxicity, useful as anti:sense molecules for treating dermatological disorders, virus infections,
              Fink JK,
                                                                        21-SEP-2001; 2001US-0323997P.
12-SEP-2002; 2002US-00242008.
                                                                                                                      13-SEP-2002; 2002WO-US029165
                                                                                                                                                                                   WO2003026566-A2
                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                           Human; atlastin; chromosome 14; 14q22.1; hereditary spastic paraplegia; HSP; neuroprotective; gene therapy; intronic splice site; gene; ds.
                                                                                                                                                                                                                                                                                                            Human atlastin exon 10 intronic acceptor splice site
                                                                                                                                                                                                                                                                                                                                          16-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                         ACF57337
                                                                                                                                                                                                                                                                                                                                                                                                     ACF57337 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 16 A; 1 C; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Fig 4a; 40pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Vasseur M, Blumenfeld M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-MAR-1993;
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                                                                                                                                                     03-APR-2003
                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16;
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              Zhao X;
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Pred. No. 5.8e+02
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Query Match
Best Local
        Matches
                                                                                                                    chromosome 14 (more specifically to 14922.1). Also described: (1) an isolared atlastin polypeptide; (2) identifying subjects who have, or are at risk of developing, hereditary spastic paraplegia (HSP); (3) a kit for determining if a subject has, or at risk of developing, HSP; (4) a computer readable medium encoding a representation of the atlastin nucleic acid sequence or polypeptide; (6) identifying subjects at risk of carrying an allele for HSP, and (7) treating a patient with HSP, Atlastin has neuroprotective activity and can be used in gene therapy. The atlastin nucleic acid is useful for preparing a composition for treating HSP or for identifying subjects who have, or at risk of developing, HSP. The present sequence represents an atlastin intronic splice site
                                                                                                                                                                                                                                                                                                                                                                                                                                                    New atlastin gene, useful for preparing a composition for treating Hereditary Spastic Paraplegia (HSP) or for identifying subjects who have, or at risk of developing, HSP.
                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                            Example 6; Page 102; 111pp; English.
                                                                                                                                                                                                                                                                                                                                                                        The present invention describes human atlastin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-371871/35.
    16;
                         Similarity
                                                                                    20 BP; 2 A; 2 C; 1 G; 15 T; 0 U; 0 Other;
        Conservative
                         1.0%;
0;
                         Score 14.2; DB 1;
Pred. No. 5.8e+02;
        Mismatches
                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                          which is located
                                            Length 20;
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Search completed: November Job time : 22 secs 2 2004, 12:44:19

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1245 ATCTTTGTTTTTTTTTA 1263

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C 1 19.6 1.4 27 1 AZB27502 C 2 18.2 1.3 24 1 AZB46788 ACCESSION.AZB27502 ACCESSION.AZB27502 ACCESSION.AZB27502 ACCESSION.AZB27502 ACCESSION.AZB26788 ACCESSION.AZB26788 ACCESSION.AZB27037 ACCESSION.AZB27037 ACCESSION.AZB27037 ACCESSION.AZB27037 ACCESSION.AZB27037 C 11 17.6 1.3 22 1 AZ318080 B 17.6 1.3 22 1 AZ319553 C 9 17.6 1.3 24 1 CF291636 ACCESSION.AZB291536 ACCESSION.AZB291737 ACCESSION.AZB291737 ACCESSION.AZB291737 ACCESSION.AZB291737 ACCESSION.AZB291747 ACCESSION.AZB291747 ACCESSION.AZB291747 ACCESSION.AZB291747 ACCESSION.AZB291747 ACC	Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution. SUMMARIES Result Query No. Score Match Length DB ID Description	gth: 8 gth: 80 inimum Match 0% aximum Match 100% leting firet 128 summar	Gapop 10- earched: 126 seqs. otal number of hits sati	Title: US-10-643-432-12 Perfect score: 1401 Sequence: 1 ccgcccgaggggacgcgagcataactaatctcttctgtt 1401 Scoring table: IDENTITY NUC	GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd. OM nucleic - nucleic search, using sw model Run on: November 2, 2004, 12:50:31; Search time 2 Seconds (without alignments) 3.826 Million cell updates/sec
C 74 15.6 1.1 22 1 CF334791 C 75 15.6 1.1 22 1 CF334791 C 75 15.6 1.1 22 1 CF334791 C 76 75 15.6 1.1 22 1 CF335250 C 77 15.6 1.1 22 1 CF335250 C 77 15.6 1.1 22 1 CF335250 C 80 15.6 1.1 22 1 CN545550 C 80 15.6 1.1 22 1 AZ316361 C 81.1 2.2 1 AZ316370 C 81.1 2.2 1 AZ316370 C 81.1 2.2 1 AZ318103 C 81.1 2.2 1 AZ424307 C 91 15.6 1.1 2.2 1 AZ424307 C 91 15.6 1.1 2.2 1 AZ459634 C 91 15.6 1.1 2.2 1 AZ463652 C 91 15.6 1.1 2.2 1 AZ696491 C 97 15.6 1.1 2.2 1 AZ7879944 C 100 15.6 1.1 2.2 1 AZ787904 C 100 15.6 1.1 2.2 1 AZ787904 C 100 15.6 1.1 2.2 1 AZ7879074 C 100 15.6 1.1 2.2 1 AZ820639 C 100 15.6 1.1 2.	65 15.6 1.1 22 1 66 15.6 1.1 22 1 67 15.6 1.1 22 1 68 15.6 1.1 22 1 70 15.6 1.1 22 1 71 15.6 1.1 22 1 72 15.6 1.1 22 1 73 15.6 1.1 22 1	56 15.8 1.1 21 1 58 15.8 1.1 22 1 58 15.8 1.1 22 1 60 15.8 1.1 22 1 61 15.8 1.1 22 1 61 15.8 1.1 22 1 62 15.6 1.1 22 1 63 15.6 1.1 22 1 64 15.6 1.1 22 1	15.8 1.1 20 1 15.8 1.1 20 1 15.8 1.1 20 1 15.8 1.1 21 1 15.8 1.1 21 1 15.8 1.1 21 1	45 15.8 1.1 19 1 46 15.8 1.1 20 1 47 15.8 1.1 20 1 49 15.8 1.1 20 1 49 15.8 1.1 20 1	. 2 1.2 22 1 1 .2 22 1 1 .2 22 1 1 .2 22 1 1 1 .2 22 1 1 1 .2 22 1 1 1 .2 22 1 1 1 .2 22 1 1 1 .2 22 1 1 1 .2 22 1 1 1 .2 22 1 1 .2 22 1 1 1 .2 22 1 1 1 .2 22 1 1 1 .2 22 1 1 1 .2 22 1 1 1 .2 23 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
ACCESSION: CF33430 ACCESSION: CF336250 ACCESSION: CF337580 ACCESSION: CF337580 ACCESSION: CF337580 ACCESSION: CS335520 ACCESSION: AZ310066 ACCESSION: AZ310066 ACCESSION: AZ311031 ACCESSION: AZ351527 ACCESSION: AZ357630 ACCESSION: AZ424307 ACCESSION: AZ424307 ACCESSION: AZ424307 ACCESSION: AZ424307 ACCESSION: AZ4263653 ACCESSION: AZ4636654 ACCESSION: AZ4636654 ACCESSION: AZ4636653 ACCESSION: AZ4636654 ACCESSION: AZ4636653 ACCESSION: AZ4636664 ACCESSION: AZ4636664 ACCESSION: AZ59664691 ACCESSION: AZ59664691 ACCESSION: AZ5966674 ACCESSION: AZ7979844 ACCESSION: AZ7987606 ACCESSION: AZ7987606 ACCESSION: AZ89841661 ACCESSION: AZ89841661 ACCESSION: AZ89841661 ACCESSION: AZ8946102 ACCESSION: AZ8946102 ACCESSION: AZ8946102 ACCESSION: AZ8946102	ACCESSION.AW332181 ACCESSION:AW332399 ACCESSION:CF299342 ACCESSION:CF31033 ACCESSION:CF311269 ACCESSION:CF311713 ACCESSION:CF311713 ACCESSION:F312498 ACCESSION:CF312498	ACCESSION: AZ610868 ACCESSION: AZ764492 ACCESSION: EX556059 ACCESSION: CO778290 ACCESSION: AZ823875 ACCESSION: AZ823875 ACCESSION: AZ823875 ACCESSION: A1708831 ACCESSION: AJ747407	ACCESSION: AZ345856 ACCESSION: AZ486784 ACCESSION: AZ588419 ACCESSION: AL048772 ACCESSION: CF318152 ACCESSION: CN546489	ACCESSION: AZ650575 ACCESSION: AL038427 ACCESSION: AL038429 ACCESSION: AL038570 ACCESSION: AL587759	ACCESSION: AL497383 ACCESSION: AX564412 ACCESSION: AZ308846 ACCESSION: AZ493766 ACCESSION: AZ493766 ACCESSION: AZ626965 ACCESSION: AZ626965 ACCESSION: AZ62734 ACCESSION: AZ601003 ACCESSION: AZ601003 ACCESSION: AZ3956341 ACCESSION: AZ3956341

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AZ827502/c
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                                                                                                                                                                                                                                                                                                                       TITLE
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123
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Insert Length: 10000 Std Error: 0.00
Plate: 0103 row: O column: 20
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                      27 bp DNA linear GSS 20-FEB-20103020R Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGC2M0103020 R, genomic survey sequence.
AZ827502
AZ827502.1 GI:12997410
GSS.
                                                                                                                                                                                                                                                                                                                     Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                              Contact: Robert B. Weiss University of Utah Genome University of Utah
                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia; 1 (bases 1 to 27)
                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus
                                                                                                                                                                                                                                                                                               Unpublished (2000)
                                                                                                                                                                                                                                                                                                             plasmid
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                                                                                                                                                                                                                                                                                                                                                                                                                            musculus (house mouse)
                                                                                                                                  quality sequence stop: 27.
Location/Qualifiers
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         /lab_host="E. Coli strain XL10-Gold, T1-resistant, P-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PW042nv; Furified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackso
Laboratory Mouse DNA Resource
                                                                                  /mor_type="genomic DNA"
/strain="C57BL/6J"
/Ab ~~~~
                                                                  /db_xref="taxon:10090"
/clone="UUGC2M0103020"
                                                                                                   /organism="Mus musculus"
/mol_type="genomic DNA"
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AL038839
CF330439
CN763587
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AZ625662
CF300172
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Sciurognathi; Muridae; Murinae; Mus.
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KEYWORDS
SOURCE
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AZ486788/c
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                                                                                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0315 row: I column: 22
Seg primer: CGTTGTAAAACGACGGCCAGT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 24)
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                                                                                                                                                                                                                                                                                                         High quality sequence stop: :
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
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University of Utah
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/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clome lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso. Laboratory Mouse DNA Resource
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                                                                                                                                                                                                     /mol_type="genomic DNA"
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                                                                                                                                                   /db_xref="taxon:10090"
/clone="UUGC1M0315122"
                                                                                                                              sex="Male"
                                                                                                                                                                                                                                                    organism="Mus musculus"
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(http://www.jax.org/resources/documents/dnares/). The DNA

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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0429 row: G column: 03
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (Dases 1 to 24)
1 (Dases 1 to 24)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., 181am,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse whole genome scaffolding with paired end reads from 10kb
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                                               /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C578L/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                /organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
(http://www.jax.org/resources/documents/dnares/). The DNI was hydrodynamically sheared by repeated passage through
                                                                                                                                                                                                                clone="UUGC1M0429G03"
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                                                                                                                                                                                                                                                                                                                                                                                               High quality sequence stop: 25.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0056 row: P column: 09
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
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Mouse whole genome scaffolding with paired end reads from 10kb
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
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                                                                                                    /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clome_lib="Mouse 10kb plasmid UUGCIM library"./note="Vector: pwp82nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
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                                                                                 Laboratory Mouse DNA Resource
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/strain="C57BL/6J"
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RESULT 5
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1520 AAAAAAAAAAAAGTAAAAGGG 1540
                                                                                                                                                                                                                                                     Barrell, Oxford Universely ....
Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available
Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20;
                                                                                                                                                                                                                                                                                                                                                                                                Constructed at the Institute for Genomic Research (TIGR), Rockville, MD. Genomic DNA isolated from a cloned population of Trypanosoma brucei (TREU927/4 GUTTAt 10.1) was mechanically shear to give a tight size distribution (4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and nhl@sanger.ac.uk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
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                                                                                                                                               /mol_type="genomic DNA"
/strain="TREU927"
/db_xref="taxon:5691"
/clone="231e08"
                                                                      1.3%;
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Pred. No. 11;
                                                                         Score 17.8;
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A clone 231e08, reverse sequence,
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CF291636
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Roslin Institute
Roslin, Midlochian, EH25 9PS, UNITED KINGDOM
Single pass sequencing. Bases called and trimmed with phred
v0.020425.c. Vector identified by cross match with the -minscore 20
and -minmatch 12 options. Vector:pBlueScriptII(KS+) R. Site 1:
ECORIR. Site 2: NotI Description: Normalised library constructed
from pooled tissue from day 30 placentas. Clones available from UK
Centre for Functional Genomics in Farm Animals, Roslin Institite,
Roslin, Midlothian, UK, EH25 9PS, www.arkgenomics.org.
                                                                                                                                                                                             Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzaee; Oryza.

1 (bases 1 to 24)
                                                                                                                                                                                                                                                                                                                                                                               sequence.
CF291636
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AJ663467 CSEQRAN09 Sus scrofa cDNA clone C0000027_007, mRNA
                                       Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                     Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus
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                                                                                     Contact: Nahm B.H
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Kyeonggi, K
31 330 6193
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/tissue_type="placenta"
/tissue_type="placenta"
/clone_Tib="CSEQRAN09"
/note="Vector: pBlueScriptII(KS+); Site_1: EcoRI; Site_2:
NotI; Single pass sequencing. Normalised library
constructed from pooled tissue from day 30 placentas."
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/db_xref="taxon:9823"
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                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0016 row: B column: 10
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus (bases 1 to 24)

Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
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Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse whole genome scaffolding with paired end reads from 10kb
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1M0016B10F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
Clone UUGC1M0016B10 F, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                           ase: plasmid ends
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                                                                                                                                                                                                                                                                                                                                quality sequence stop: 24.
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/cultivar="Nackdong"
/db_xref="taxon:39947"
/db_xref="taxon:39947"
/clone="14ROOT--02-C09"
/tlesue_type="root"
/dev_stage="14 days after germination"
/dev_stage="14 days after germination"
/lab_host="E.coli DH10B"
/lab_host="E.coli DH10B"
/clone_lib="Rice_root plasmid_cDNA_library_(14ROOT)"
/clone_lib="Rice_root plasmid_cDNA_library_(14ROOT)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA_was capped
with oligoribonucleotides and then used as templates for
RT-PCR."
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                   /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PWD42nv, Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biomedical Polymers Research Bldg., 20
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http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                db_xref="taxon:10090"
clone="UUGC1M0016B10"
                                                                                                                                                                                                          mol_type="genomic DNA"
etrain="C57BL/6J"
                                                                                                                                                                                                                                                           organism="Mus musculus"
                                                                                                                                         sex="Male"
                                                                                                                                                                                                                                                                                                            ocation/Qualifiers
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83.3%;
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Pred. No. 16;
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as templates for
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TITLE JOURNAL COMMENT

REFERENCE

AUTHORS

FEATURES

Query Match Best Local

1.3%; 1 Similarity 83.3%; 20; Conservation

Pred. No. 17; 0; Mismatches

DB 1;

Length 25;

4.

Indels

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Gaps

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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeenggi, Korea Tel: 82 31 330 6193

Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Cryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae; Streptophyta; Embryophyt
Spermatophyta; Magnoliophyta; Liliopsida; Poales;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25 pp mRNA linear EST 15-AUG-2003 HD--06-I14.gl OsHDAC1-overexpressing transgenic rice plasmid cDNA library (HD) Oryza sativa (japonica cultivar-group) cDNA clone HD--06-I14, mRNA sequence.
CF317007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Large-scale Sequencing Analysis Unpublished (2003) Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr
/note="Vector: pCR4-TOPO; Site 1: EcoRI; Callus was treated with ABA(20um) for 1hr. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was derived from rice Histone Deacetylase overexpression line."
                                                                                                                                       was hydrodynamically sheared by repeated passage through
                                                                                                                                                                                                                                                                                                                                     /organism="Oryza sativa
/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                    tissue_type="callus"
                                                                                                                                                                                                                                                                              /db_xref="taxon:39947"
/clone="HD--06-I14"
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Pred. No. 16
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                                                                                                                                                                                                                                                                                                                                                                                           (japonica cultivar-group)"
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ACCESSION VERSION KEYWORDS

SOURCE

ORGANISM

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Matches

POCUS

DEFINITION CF317007/c RESULT 9

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                  RESULT 11
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JOURNAL
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                                                                                                                                                                                                 Local Similarity
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                                                                                                  24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: -21m13
High quality seque
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           High qality sequence starts: 1
High qality sequence stops: 1
Source: IMAGE Consortium, LINL
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Generation and analysis of 280,000 human expressed sequence tags
Genome Res. 6 (9), 807-828 (1996)
97044478
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  AZ307896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Other_ESTs: yb08h08.rl
Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and Marra, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T49097.1 GI:650957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TTAATTAAAAAAAAAAAAGTAAAA 1537
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314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          quality sequence stop: 1.
                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="GDB:491520"
                                                                                                                                                                                                                                                                                                                                                                                                                     /db_xref="taxon:9606"
/clone="IMAGE:70623"
                                                                                                                                                                                                                                                                                                                                                                                                  sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ocation/Qualifiers
                                                                                                                                                                                               1.3%;
                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                             Score 17.6;
Pred. No. 17
  22
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  DNA
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  linear
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  GSS 29-SEP-2000
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                      RESULT 12
CF308058/c
LOCUS
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JOURNAL COMMENT
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AUTHORS
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                                                                                                                                          Query Match
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                                                                                                                    Local
                                             1520 AAAAAAAAAAAGTAAAAGGGA 1541
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erry
Plate: 0010 row: N column: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Welss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1M0010N18F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0010N18 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      High quality sequence stop:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dunn, D., Aoyagi, A., Barber, M.,
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AZ307896.1
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                                                                                             Conservative
                                                                                                                                                                                http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNA42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'sex="Male'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mol_type="genomic DNA"
strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     clone="UUGC1M0010N18"
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                                                                                                                    1.2%;
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                                                                                                                                        Score 17.2;
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                                                                                             Mismatches
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                                                                                                                                             DB 1;
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CF308058 23 bp mRNA linear collistance ABF--01-L15.gl ABF3-overexpressing transgenic rice plasmid cDNA

EST 15-AUG-2003

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JOURNAL COMMENT
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AUTHORS
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AZ627850
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AUTHORS
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Oryza sativa (japonica cultivar-group)
Cryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta; Tracheophyta;
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae;
Ehrhartoideae, Oryzeae, Oryza.
plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                        Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 23) Kim, M.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Shin, Y.C., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                            Mus musculus
                                                                                                                                                                                                                                                                                                                                    AZ627850.1 GI:11750136 GSS.
                                                                                                                                                                                                                                                                                                                                                                                            Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea
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CF308058
CF308058.1 GI:33679819
                                                                                                                                                                                                                                                                                                               Mus musculus (house mouse)
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 library (ABF) Oryza sativa (japonica cultivar-group) cDNA clone
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="Vector: pCR4-TOPO; Site 1: ECORI; Leaf was dried for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /dev stages"14 days after germination"
/lab_host="E.coli DH108"
/clone lib="ABF3-overexpressing transgenic rice
cDNA library (ABF)"
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Pred. No. 20;
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RESULT 14
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CF300172 23 bp mRNA linear EST 15-AUG-: 7LEAF--04-H15.b1 Rice leaf plasmid cDNA library II (7LEAF) Oryz: sativa (japonica cultivar-group) cDNA clone 7LEAF--04-H15, mRNA
                                                                                                                                            Kim,J.S., Jun,K.M., Cheong,P.J.,
Song,S.I., Kim,J.K., Kim,Y.-K. an
Large-scale Sequencing Analysis o
Unpublished (2003)
of Bioscience and Bioinformatics,
Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Insert Length: 10000 Std Err
Plate: 0474 row: N Column: 2
                                                                      Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                  Contact: Nahm B.H.
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Fax: 801 585 7177
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                                                                                                                                                                                                                                               (bases 1 to 23)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XI10-Gold (Stratagene) cells
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and selected for ampicillin resistance."
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/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="genomic DNA"
/strain="C57BL/6J"
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. and Nahm,B.H.
is of Rice ESTs
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ACCESSION
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JOURNAL
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AL038576
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AUTHORS
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VERSION
KEYWORDS
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Best Local Similarity
Matches 18; Conserv
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Best Local Similarity
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           AL038576 20 by DKFZp566F0346 rl 566 (synonym: DKFZp566F0346, mRNA sequence. AL038576
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DKFZp566D1846 rl 566 (synonym: hfkd2) Homo
DKFZp566D1846, mRNA sequence.
AL038507
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Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 20)
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 AL038576.1
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ottenwaelder, B., Obermaier, B., Mewes, H.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Wiemann,S.
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ilarity 90.0%;
Conservative
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/clone="DKEZD56611846"
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/dev_stage="fetal"
/lab_host="x1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI; Site_2: Sa
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /cullivar="Nackdong"
/cb xref="Nackdong"
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/clone lib="Rice leaf plasmid cDNA library II (7LEAF)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped with oligoribonucleotides and then used as templates for RT-pCR."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'organism="Oryza sativa (japonica cultivar-group)"
'mol_type="mRNA"
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/ 100.0%; Pr/
GI:49682161
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Pred. No.
                                                                                                                                                                                      Score 16.8;
Pred. No. 2:
                                                                                                                                                                          Mismatches
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                                                   20 bp
                                       hfkd2) Homo
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sapiens
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                                       sapiens
                                                     linear
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                                       EST 06-JUL-2004
DNA clone
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JOURNAL
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AL038754
LOCUS
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AUTHORS
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Best Local S
Matches 18
                                                      Matches
                                                                 Query Match
Best Local
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                                                      18;
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18; Conservative
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Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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EST.
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1. (bases 1 to 20)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DKFZp566M1346 rl 566 (synonym: DKFZp566M1346, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EST (Ottenwaelder, et al.)
Unpublished (1999)
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Ottenwaelder, B., Obermaier, B.,
                                                                                                                                                                                                                                                                                                    Ingolstaedter Landstr.1, D-85764 Neuherberg, Germany.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                               Ottenwaelder,B., Obermaier,B.,
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    Location/Qualifiers
TTAAAAAAAAAAAGTAAAA 1537
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                                                                                                                                                                                          /organism="Homo sapiens"
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/db_xref="taxon:9606"
/clone="pKFZp566M1346"
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/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: Not
                                                                                                                   /note="Vector: pAMP1; Site_1: NotI; Site_2:
                                                                                                                                       /dev_stāge="fetal"
/lab_host="Xl-2blue"
/clone_lib="566 (synonym: hfkd2)"
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/mol_type="mRNA"
/db_xref="taxon:9606"
/clone="DKFZp566F0346"
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90.0%; Pred. No. 22;
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                                                  0; Mismatches
                                                                   Score 16.8;
Pred. No. 2;
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hfkd2) Homo
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                                                                                   DB 1;
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sapiens
                                                    Indels
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cDNA clone
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VERSION
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CF298207
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AUTHORS
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VERSION
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AL038845
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Best Local S
Matches 18
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JOURNAL
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                                                                                                                                                                                                                   Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.

1 (bases 1 to 20)
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CF298207

CF298207

7LEAF--01-H23.gl Rice leaf plasmid cDNA library II (7LEAF) Oryza sativa (japonica cultivar-group) cDNA clone 7LEAF--01-H23, mRNA
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1 (Dases 1 to 20)

Ottenwaelder, B., Obermaier, B., Mewes, H.W., Gassenhuber, J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DKFZp566P1746 rl 566 (synonym: DKFZp566P1746, mRNA sequence.
                                                                                                   Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                            EST
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Unpublished (1999)
                                                                                                                                                             Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                CF298207.1 GI:33669968
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Location/Qualifiers
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                                                                                                                                                                                                      Contact: Nahm B.H.
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                                                            bhnahm@ggbio.com, bhna
Location/Qualifiers
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/db xref="taxon:9606"
/db xref="taxon:9606"
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/dev stage="fetal"
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/note="Vector: pamP1; Site_1: NotI; Site_2:
/organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      organism="Homo sapiens"
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                                                                                                                                            Korea
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Pred. No. 22
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                                                                                  bhnahm@bio.myongji.ac.kr
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nym: hfkd2) Homo sapiens cDNA clone
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0150 row: P column: 21
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Mouse whole genome scaffolding with paired end reads from 10kb
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1. (bases 1 to 20)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert
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Unpublished (2000)
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
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/db xref="taxon:39947"
/clone="JLEAF-01-H23"
/tissue_type="leaf"
/dev_stage="7 days after germination"
/leb_host="E.coli DHLOB"
/clone_lib="Rice leaf plasmid cDNA library II (7LEAF)"
/clone_lib-"Rice leaf plasmid cDNA library II (7LEAF)"
/clone_library II (7LEAF)"
                                                                                                                                                                                                                                                                                                                       Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone_lib="Mouse_10kb_plasmid_UUGCiM_library" /note="Vector: pW042nv; Purified_genomic_DNA_from_M. musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_from_the_Jackson_was_obtained_fro
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/strain="C57BL/6J"
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Pred. No. 22;
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smid UUGC1M library Mus musculus genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ddurn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0203 row: K column: 21
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (2000)
Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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ZMO203K21R Mouse 10kb plasmid UUGC2M library Mus musculus genomic
clone UUGC2M0203K21 R, genomic survey sequence.
AZ943299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 High quality sequence stop: 21.
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mouse whole genome scaffolding with paired end reads from 10kb
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                          (http://www.jax.org/resources/documents/dhares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                /lab host="E. coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse 10kb plasmid UUGC2M library" /note="Vector: PWD42nv; Purified genomic DNA from M.musculus C57BL/6J (female) was obtained from the Jacks
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
10.5 kb range using preparative agarose gel
                                                                                                                                                                                                                                 Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:10090"
/clone="UUGC2M0203K21"
                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                           sex="Female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  organism="Mus musculus"
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90.0%;
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Pred. No. 22;
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COMMENT

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KEYWORDS
                                                                                                                                                                                                                                 RESULT 23
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                                       ORGANISM
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                                                                                                                                                                                                                                                                                                                                                                     18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21 AAAAAAAAAAAAAAAAGGG 2
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                               CF310486 22 bp mRNA linear EST 15-AUG-2003 ABF--05-C16.gl ABF3-overexpressing transgenic rice plasmid cDNA library (ABF) Oryza sativa (japonica cultivar-group) cDNA clone ABF--05-C16, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DKFZp566C1646_r1 566 (synonym:
                                                                             EST.
                                                                                             CF310486
CF310486.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EST (Ottenwaelder, et al.)
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wiemann, S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 22)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: MIPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ottenwaelder, B., Obermaier, B., Mewes, H.W., Gassenhuber, J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ingolstaedter Landstr.1, D-85764 Neuherberg, Germany
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      electrophoresis. Vector DNA was prepared from a derivative of pWp42 (gi|4732114|gb|AFI19072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI; Site_2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="DKFZp566C1646"
/tissue_type="kidney"
/dev_stage="fetal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ocation/Qualifiers
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Pred. No. 25;
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hfkd2) Homo
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CF329694/c
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., L. Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                 Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta;
Spermatophyta, Magnoliophyta, Liliopsida, Foales, Poaceae,
Ehrhartoideae, Oryzeae, Oryza.
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NACL--05-B19.b1 Rice callus plasmid cDNA library (NACL) Oryza
sativa (japonica cultivar-group) cDNA clone NACL--05-B19, mRNA
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Shin, Y.C.,
Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                         Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
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EST.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Nahm B.H
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                                                                                                                                                                                                                                                                     bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr.
Location/Qualifiers
/cultivar="Nackdong"
/db_xref="taxon:39947"
/clone="NACL--05-B19"
/tissue_type="callus"
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/lab_hosts"E.coll DH10B"
/clone libs"ABF3-overexpressing transgenic rice plasmid
/clone library (ABP)"
/clone Type College Coll
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/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                       /organism="Oryza sativa
/mol_type="mRNA"
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/clone="ABF--05-C16"
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Pred. No. 25;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shin, Y.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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CF333801/c
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ORGANISM
                     DEFINITION
                                                                          RESULT 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Nahm B.H.
Genomics and Genetics Institute, GreenGene Biotech In of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeenggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnollophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JMT--02-N11.g1 AtJMT-overexpressing transgenic rice plasmid cDNA library (JMT) Oryza sativa (japonica cultivar-group) cDNA clone JMT--02-N11, mXNA sequence.
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CF333801.1 GI:33815910
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: bhnahm@ggbio.com,
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                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                   was reverse transcribed and then used for PCR. mRNA was pepared from Arabidopsis Jasmoinate Carboxyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="Rice callus plasmid cDNA library (NACL)" /note="Vector: pCR4-TOPO; Site_l: EcoRI; mRNA was capped with oligoribonuclectides and then used as templates for
                                                                                                                                                                                                                                                                                   methyltransferase overexpression
                                                                                                                                                                                                                                                                                                                                                        /clone lib="AtJMT-overexpressing transgenic rice plasmid cDNA library (JMT)"
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                                                                                                                                                                                                                                                                                                                                          note="Vector: pCR4-TOPO; Site_1: EcoRI; Oligo-capped mRNA
                                                                                                                                                                                                                                                                                                                                                                                               /dev_stage="14 days after germination"
/lab_host="E.coli DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                  tissue_type="leaf"
                                                                                                                                                                                                                                                                                                                                                                                                                                                       'db_xref="taxon:39947"
'clone="JMT~-02-N11"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cultivar="Nackdong"
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mol_type="mRNA"
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lab_host="B.coli DH10B"
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90.0%; Pred. No. 26
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Pred. No. 26;
0; Mismatches
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                   C0000045_P10, mRNA
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RESULT 27
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Best Local
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                                                                                                                                                                                                                                                                                                                      Glossina morsitans morsitans Glossina morsitans (Glossina morsitans morsitans morsitans morsitans morsitans morsitans Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Hippoboscoidea; Glossinidae; Glossina.

1 (bases 1 to 23)
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Anderson, S.I., Finlayson, H.A. and Archibald, A.L.
Development of cDNA and BST resources for studying reproduction embryo development in pigs and cattle
Unpublished (2004)
                                    Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 ISA, UK
Request for clones, please contact: Mike Lehane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23 bp mRNA linear EST 10-C BX550903 Glossina morsitans morsitans adult infected gut Glo morsitans morsitans morsitans adult infected gut Glo morsitans morsitans cDNA clone Tsell5e01_qlc, mRNA sequence. BX550903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Roslin Institute
Roslin, Midlothian, EH25 9PS, UNITED KINGDOM
Single pass sequencing. Bases called and trimmed with phred
Single pass sequencing. Bases called and trimmed with phred
v0.020425.c. Vector identified by cross match with the -minscore 20
and -minmatch 12 options. Vector:pBlueSCriptII(KS+) R. Site 1:
ECORI R. Site 2: NotI Description: Normalised library constructed
from pooled tissue from day 30 placentas. Clones available from UK
Centre for Functional Genomics in Farm Animals, Roslin Institute,
Roslin, Midlothian, UK, EH25 9PS, www.arkgenomics.org.
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Genomics and Bioinformatics
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EST.
                                                                                                                               14519198
Contact: Hall N
                                                                                                                                                                                             response genes
Genome Biol. 4
                                                                                                                                                                                                                                        morsitans morsitans and expression analysis of putative immune
                                                                                                                                                                                                                                                                                   Lehane,M.J., Aksoy,S., Gibson,W., Kerhornou,A., Be
Hamilton,J., Soares,M.B., Bonaldo,M.F., Lehane,S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BX550903.1 GI:33374645
                                                                                                                                                                           22881942
                                                                                                                                                                                                                                                                 Adult midgut expressed sequence tags from the tsetse fly Glossina
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Biological Sciences,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           organism="Sus scrofa"
/mol_type="mRNA"
/db_xref="taxon:9823"
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Pred. No. 28
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S. and Hall,
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REFERENCE
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                                                                                                                                                                                                                                                                                                                                                         Contact: Hall N
Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for clones, please contact: Mike Lehane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Glossina morsitans morsitans
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Hippoboscoidea; Glossinidae; Glossina.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23 bp mRNA linear EST 14-
BX568055 Glossina morsitans morsitans adult infected gut Gl
morsitans morsitans cDNA clone Tse91f03_plc, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  University of Wales,
Bangor LL57 2UW
All clones with suffix q1c are reverse
end of the cDNA all plc reads are from
the 3' end.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lehane,M.J., Aksoy,S., Gibson,W., Kerhornou,A., Berriman,M., Hamilton,J., Soares,M.B., Bonaldo,M.F., Lehane,S. and Hall,N. Adult midgut expressed sequence tags from the testes fly Glossina morsitans morsitans and expression analysis of putative immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BX568055
                                                                                                                                                                                                                                                                                   University of Wales,
Bangor LL57 2UW
                                                                                                                                                                                                                                                                                                                           Prof. M.J.Lehane
School of Biological Sciences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     response genes
Genome Biol. 4 (10), R63 (2003)
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                                   /clone="Tse91f03_p1c"
/tissue_type="adult infected gut"
/clone_lib="Glossina morsitans mo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone="Tsel15e01 q1c"
/tissue_type="adult infected gut"
/clone_lib="Glossina morsitans morsitans adult infected
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|mol_type="mRNA"
|sub_speciese="morsitans"
|db_xref="taxon:37546"
                                                                                                                   sub_species="morsitans"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           note="country: Zimbabwe; EST from adult gut infected with
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                                                                                                                                      mol_type="mRNA"
                                                                                                                                                           organism="Glossina morsitans morsitans"
                                                                                                                                                                                                   ocation/Qualifiers
note="country: Zimbabwe; EST from adult
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GI:33434952
                                                                                                    xref="taxon:37546"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Score 16.6; D; Pred. No. 30; 0; Mismatches
                                                                                                                                                                                                                                             ereverse
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AZ486853/c
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AZ486853 GI:10654033
AZ486853.1 GI:10654033
GSS.
Mus musculus (house mouse
Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop: 23.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0315 row: M column: 14
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
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Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T.,
Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von
Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 23)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert B.
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pund2 (gi[473214]gb]AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      /lab host="E. Coli strain XL10-Gold, T1-resistant, F-" /Clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: pWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                   Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mol_type="genomic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             clone="UUGC1M0315M14"
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Pred. No. 30;
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ORGANISM
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1M0510005R Mouse 10kb plasmid UUGCIM library Mus musculus genomic
clone UUGC1M0510005 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0510 row: O column: 05
Seg primer: CACACAGGAAACAGCTATGACC
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University of Utah Genome Center
University of Utah
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Location/Qualifiers
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Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von
Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AZ645254.1 GI:11774572
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               (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pDNA2 (gi|4732114|gb|AF1290721), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: pW042nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
  and selected
                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               clone="UUGC1M0510005"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 strain="C57BL/6J"
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Pred. No. 30;
for ampicillin resistance."
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Pred. No. 30;

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REFERENCE
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19 bp DNA line 1M0529F08F Mouse 10kb plasmid UUGCIM library Muclone UUGCIM0529F08 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0529 row: F column: 08
Seq primer: CGTTGTAAAACGACGGCCAGT
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Fax: 801 585 7177
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Mammalla; Eutheria; Rodentia;
1 (Dases 1 to 19)
Dunn, D., Aoyagi, A., Barber, M.,
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Location/Qualifiers
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10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                      (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 6 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                     /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone_lib="Mouse 10kb plasmid UUGCIM library. Colone_lib="Mouse 10kb plasmid UUGCIM library. Incre="Vector: pwp42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso
                                                                                                                                                                                                                                                                                                                                                                                                                                  Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="genomic DNA"
/strain="C57BL/6J"
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Sciurognathi; Muridae; Murinae; Mus
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Query Match Best Local Similarity

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Matches Query Match Best Local (

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Nonce whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0077 row: E column: 20
Seq primer: CGTTGTAAAACGACGGCCAGT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert B. Weiss
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|AF12972.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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T. brucei sheared genomic Di
genomic survey sequence.
AL497383 1 GI:11865504
GSS.
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Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UX
Request for Clones, please contact: Mike Lehane
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Trypanosoma brucei
Trypanosoma brucei
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Prof. M.J.Lehane
School of Biological Sciences,
University of Wales,
Bangor LL57 2UW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lehane, M.J., Aksoy, S., Gibson, W., Kerhornou, A., Berriman, M., Hamilton, J., Soares, M.B., Bonaldo, M.F., Lehane, S. and Hall, N. Adult midgut expressed sequence tags from the tsetse fly Glossina morsitans morsitans and expression analysis of putative immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Hippoboscoidea; Glossinidae; Glossina.
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of the cDNA all p1c reads are from
3' end.
                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                   /note="country: Zimbabwe; T.brucei"
                                                                                                                                                                                                                                                                                                                                                                                                                  /clone="Tse101g03_p1c"
/tissue_type="adult infected gut"
/clone_Tib="Glossina morsitans mo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Glossina morsitans morsitans"
/mol type="mRNA"
/sub_species="morsitans"
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A clone
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                                       Prof. M.J.Lehane
School of Biological Sciences,
University of Wales,
Bangor LL57 2UW
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BX564412 Glossina morsitans morsitans adult infected gut Glossina morsitans morsitans morsitans morsitans clone Tse7le10_plc, mRNA sequence.
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4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
All clones with suffix qlc are reverse primer reads starting at end of the cDNA all plc reads are from the 3' end.
                                                                                                                                                       Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for clones, please contact: Mike Lehane
                                                                                                                                                                                                                                                                                                                                 response genes
Genome Biol. 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Hippoboscoidea; Glossinidae; Glossina.
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Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nchl@sanger.ac.uk
Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.B., Rajandream,M.A. and Barrell,B.G.
                                                                                                                                                                                                                                                           Contact: Hall N
                                                                                                                                                                                                                                                                                      14519198
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/strain="TREU927"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0012 row: H column: 15
Seg primer: CGTTGTAAAACGACGGCCAGT
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Mus musculus (house mous
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1 (bases 1 to 21)
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1M0012H15F Mouse 10kb plasmid UUGCIM library Mus musculus genomic
clone UUGC1M0012H15 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Class: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      308,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                     quality sequence stop: 21.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
                  (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4
                                                                                                                               /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso
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polynucleotide kinase. Adaptor oligonucleotides were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone="Tse71e10 p1c"
/tissue_type="adult infected gut"
/clone_lib="Glossina morsitans morsitans adult infected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                 Laboratory Mouse DNA Resource
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/db_xref="taxon:37546"
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                                                                                                                                                                                                                                                     /db_xref="taxon:10090"
/clone="UUGC1M0012H15"
                                                                                                                                                                                                                                                                                              /mol_type="genomic DNA"
                                                                                                                                                                                                                                                                                                                                             organism="Mus musculus"
                                                                                                                                                                                                                                   sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note="country: Zimbabwe; EST from adult gut infected with
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Weiss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 16.2;
Pred. No. 35;
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                                                                                                                                      from the Jackson
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                                                                                            The DNA
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AUTHORS
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AZ493766
LOCUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURES
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ORGANISM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0328 row: C column: 11
Seg primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 21)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AZ493766 21 bp DNA linear GSS 05-OCT-200 IM0328C11R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0328C11 R, genomic survey sequence.

AZ493766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Class: plasmid ends
High quality sequence stop: 21.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AZ493766.1 GI:10667750
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 6 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                           /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clome_lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWB42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso
                                                                                                                                                                                                                                                                                                                                                                                          /organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                    Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                            /clone="UUGC1M0328C11"
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Pred. No. 35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Std Error: 0.00
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RESULT 38
AZ513902/c
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Best Local Similarity
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Rose,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
Nouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seq primer: CGTTGTAAAACGACGGCCAGT Class: plasmid ends High quality sequence stop: 21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0360 row: A column: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert B.
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GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (2000)
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(http://www.jax.org/resources/documents/dhares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                          /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-/clone_lib="Mouse_10kb_plasmid_UUGCLM library" /note="Vector: PWD42nv, Purified_genomic_DNA_from M.musculus_C57BL/6J_(male) was obtained from the Jack Laboratory_Mouse_DNA_Resource_
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/strain="C57BL/6J"
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clone="UUGC1M0360A13"
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Pred. No. 35
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REFERENCE AUTHORS

COMMENT

JOURNAL

FEATURES

VERSION KEYWORDS

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REFERENCE
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Matches 18
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,B., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0467 row: E column: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          plasmid inserts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (house mouse)
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Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel
                                                                                                                                                                                                                                                                                                                                                                                                /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /Clone_lib="Mouse_10kb_plasmid_UUGCIM_library" /note="Vector: PW042nv, Purified_genomic_DNA_from_M.musculus_C57BL/6J_(male) was obtained_from_the_Jackson_musculus_C57BL/6J_(male) was obtained_from_the_Jackson_Musculus_C57BL/6J_(mal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            db xref="taxon:10090"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          clone="UUGC1M0467E15"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Trace considered overall poor quality
Insert Length: 520 Std Error: 0.00
Seq primer: -40m13 fwd. ET from Amersham
High quality sequence stop: 1.
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AA911173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AA911173 22 bp mRNA linear EST 09-JUN-19: ok81a10.s1 NCI_CGAP_Kid3 Homo sapiens cDNA clone IMAGE:1520346 3' similar to TR:\(\overline{Q}34192\) Q34192 NADH DEHYDROGENASE SUBUNIT 5. ;, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tumor Gene Index
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA Library Preparation: M. Bento Soares, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
cound through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TGTTGTGTGTGTGTGCCTG 21
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                                                                                         /clone lib="NCI CGAP_Kid3"
/clone lib="NCI CGAP_Kid3"
/note="Organ: kidney; Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Ecc RI; 1st strand cDNA was primed with a Not I - oligo(dT) primer, double-stranded cDNA was ligated to Ecc RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Ecc RI sites of the modified pT7T3 vector: mENA source: 2 pooled kidneys. Library went through one round of normalization. Library constructed by Bento Soares and M. Fatima Bonaldo. "
                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:9606"
/clone="IMAGE:1520346"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                organism="Homo sapiens"

mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ocation/Qualifiers
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85.7%;
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Pred. No. 37
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Insert Length: 10000 Std Error: 0.00
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1M0542D04F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0542D04 F, genomic survey sequence.
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Fax: 801 585 7177
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Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                    was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymicleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of phasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/strain="C57BL/6J"
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Location/Qualifiers
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Insert Length: 10000 Std Error: 0.00
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Fax: 801 585 7177
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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/strain="C57BL/6J"
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2M0222H16R Mouse 10kb plasmid UUGC2M library Mus musculus genomic
clone UUGC2M0222H16 R, genomic survey secuence
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84112, USA
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Seg primer: CACACAGGAAACAGCTATGACC
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    University of Utah Genome Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2000)
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                            Similarity
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                                                                                                      adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored me user adaptors and purified vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                             (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /lab host="E. coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGC2M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
/musculus C57BL/6J (female) was obtained from the Jackson
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/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             organism="Mus musculus"/
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  100.0%; F
tive 0;
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Score 16; DB 1; Length 21; pred. No. 40; 0; Mismatches 0; Indels
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AZ345795
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                                                                                                 l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              High quality sequence stop: 19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0080 row: H column: 09
Seq primer: CACACAGGAAACAGCTATGACC
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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AZ345795.1 GI:10425032
                                       TTAAAAAAAAAAAGTAAA 1536
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801 585 7177
                                                                                                                                                                                                                                                                (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pubd2 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector NNA and transformed into
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                                                                                                                                                                                                   adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
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/strain="C57BL/6J"
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                                                                                                                         1.1%;
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                                                                                                                            Score 15.8;
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                                                                                                                                                 DB 1;
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REFERENCE
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Nouse whole genome scaffolding with paired end reads from 10kb mouse whole genome scaffolding with paired end reads from 10kb
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Insert Length: 10000 Std Error: C
Plate: 0520 row: P column: 13
Seq primer: CACACAGGAAACAGCTATGACC
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Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Location/Qualifiers
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Fax: 801 585 7177
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                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi 4732114 |gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/strain="C57BL/6J"
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1.1%;
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smid UUGCIM library Mus musculus genomic
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VERSION
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AL038427
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Best Local Similarity
Matches 17; Conser
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Unpublished (1999)
                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1. (bases 1 to 20)
                                                                                                                                                                                                                                                                                                                                                                                                                           DKFZp566A1946_rl 566 (synonym: DKFZp566A1946, mRNA sequence.
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                                                                                                                                                                                                                                                                                Ottenwaelder, B., Obermaier, B.,
                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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Location/Qualifiers
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Unpublished (1999)
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Mammalia; Eutheria; Primates;
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Location/Qualifiers
                                                                                                                                                                                                                 Contact: MIPS
                                                                                                                                                                                                                                                                   Wiemann, S
                                                                                                                                                                                                                                                                                                                                                               Homo sapiens (human)
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/organism="Homo sapiens"
/mol type="mRNA"
/db xref="taxon:9606"
/clone="DKRZD566A1946"
/tlssue_type="kidney"
/dev_stage="fetal"
/lab hoste="X1-Zblue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI; Si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: Not
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/tissue_type="kidney"
/dev_stage="fetal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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89.5%;
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Pred. No. 43
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nym: hfkd2) Homo
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hfkd2) Homo sapiens cDNA clone
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      Site_2: SalI"
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AL587759/c
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Best Local
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Dept. Genom:
Roslin Institute
Roslin, Midlothian, EP
Rosl: +44 (0)131 527 4;
Fax: +44 (0)131 440 0
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                                                                                                                                                                               Gallus gallus
                                                                                                                                                                                                                                          AL587759 20 bp
AL587759 BP Chicken Brain Library
ROS061G06, mRNA sequence.
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                                                                          Contact: Frazer Murray
                                                                                        Unpublished (2001
                                                                                                   Murray,F.
BP Chicken Brain Library
                                                                                                                                                                                            Gallus gallus (chicken)
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Location/Qualifiers
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Unpublished (1999)
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Ottenwaelder, B., Obermaier, B.,
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Mammalia; Eutheria;
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                                                              Genomics and Bioinformatics
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/db_xref="taxon:9606"
/clone="DKFZp566F1746"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /tissue_type="kidney"
/dev_stage="fetal"
/lab_host="X1-2blue"
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             EH25
4200
0434
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Primates;
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0; Mismatches
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Pred. No. 43
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Pred. No. 43;
                                     9PS, UK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 20)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
Nouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error: 0.00
Plate: 0080 row: G column: 17
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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University of Utah Genome Center
University of Utah
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AZ345856.1 GI:10425093
GSS.
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Fax: 801 585 7177
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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GCGGCCGCTTTTTTTTTTTTTTTTTTTT 3' Poly A RNA purchased from Clonetech
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TAAAAAAAAAAAAGTAAAA 1537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 308,
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                                                                                                                                                                                                                                                                                                                                                                                                            quality sequence stop: 20.
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/note="Vector: pSPORT1; Site 1: Not1; Site 2: Sal1; Cloned
/note="Vector: pSPORT1; Site 1: Not1; Site 2: Sal1; Cloned
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/dev_stage="Unknown"
/lab_host="DH10B"
/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clome_lib="Mouse 10kb plasmid UUGC1M library"
/note="Yector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                 mol_type="genomic DNA"
strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                  organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                  ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /db_xref="taxon:9031"
/clone="ROS061G06"
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|mol_type="mRNA"
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                                                                                                                  sex="Male"
                                                                                                                                                    clone="UUGC1M0080G17"
                                                                                                                                                                                       db_xref="taxon:10090"
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Pred. No. 43;
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UUGC1M library Mus musculus
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AZ486784/c
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weise,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AZ486784 20 bp DNA linear GSS 0:
1M0315C20F Mouse 10kb plasmid UUGC1M library Mus musculus
clone UUGC1M0315C20 F, genomic survey sequence.
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17; Conserv
                                                                                                                                                                                                                                                                                                   Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 20
                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0315 row: C column: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       University of Utah Genome Center University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
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/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Wecttor: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                          /db_xref="taxon:10090"
/clone="UUGC1M0315C20"
                                                                                                                                                                           /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                             organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GI:10653898
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Pred. No. 4:
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KEYWORDS

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AZ858419
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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2M0163003R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0163003 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seq primer: CACACAGGAAACAGCTATGACC Class: plasmid ends High quality sequence stop: 20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0163 row: O column: 03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNAPA2 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and sheared adaptors on the sheared adaptors of the sheared sheared mouse nna annealed to the sheared with adaptors and sheared adaptors of the sheared sheared
/lab_host="E. Coli strain XL10-Gold, T1-resistant, F /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jack Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Biomedical Polymers Research Bldg., 20 S. 2030 E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                      clone="UUGC2M0163003"
                                                                                                                                                                                                                                                                                                                               /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                     organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocation/Qualifiers
                                                                                                                                                                                                               sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.1%;
                                                                                                                                                                                                                                                                                        _xref="taxon:10090"
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Pred. No. 43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Std Error: 0.00
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                                                          RESULT 54
CF318152/c
LOCUS
DEFINITION
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DEFINITION
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AL048772
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Best Local S
Matches 17
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Best Local Similarity
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                                                                                                                                                                                                                   1513 GTTAATTAAAAAAAAAAA 1531
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                                   CF318152

10-08-C11.bl OsHDAC1-overexpressing transgenic rice plasmid of library (HD) Oryza sativa (japonica cultivar-group) cDNA clone
                                                                                                                                                                                                                                                               17;
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DKFZp566N143 rl 566 (synonym:
DKFZp566N143, mRNA sequence.
                                                                                                                                                                                N
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HD--08-C11, mRNA sequence. CF318152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 21)
Kochter, K., Beyer, A., Mewes, H.W.,
EST (Kochter, et al.)
Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ingolstaedter Landstr.1, D-85764 Neuherberg, Germany.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: MIPS
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                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                GTTAAAAAAAAAAAAAA 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens (human)
                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide Kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                          /db xref="taxon:9606"
/clone="bKFZp566N143"
/tissue="pe="kidney"
/tissue_tpe="fetal"
/dev stage="fetal"
/lab_host="x1-2blue"
/clone_lib="566 (synonym: hfkd2)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                                                                                                                                                                                                        note="vector: pamp1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GI:4727843
                                                                                                                                                                                                                                                                                1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   _type="mRNA"
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                                                                                                                                                                                                                                                                                Pred.
                                                                                                                                                                                                                                                                                  Score 15.8;
Pred. No. 46;
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Pred. No. 4
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: hfkd2) Homo sapiens cDNA clone
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                                                                                                                                                                                                                                                                                                  DB 1;
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                                                                                                                                                                                                                                                                                                  Length 21;
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                                                                                 EST 15-AUG-2003
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CN546489/c
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AUTHORS
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                                                                                                                                 COMMENT
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Best Local (
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                                                                                                                                                                                                 BTILE
                                                                                                                                                                                                                                                                    AUTHORS
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                                                                                                                                                     JOURNAL
                                                                                                                                                                                                                                                                                                                                                                              ORGANISM
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17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                        Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; rosids; Vitaceae; Vitis.

1 (bases 1 to 21)
1 (bases 1 to 21)
Abbal, P., Agasse, A., Ageorges, A., Atanassova, R., Barrieu, F., Abbal, P., Dedaldechamp, F., Delrot, S., Glissant, D., Grimplet, J., Hamdi, S., Romieu, C. and Terrier, N.
Generation of Expressed Sequence Tag from Grape Berry (skin, pulp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CF318152.1 GI:33689913
EST.
                                                                                                   or seeds) at Various Developmental Stages
Unpublished (2002)
Contact: Hamdi S.
UMR 619 - Equipe Biologie de la Vigne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EST 18633 Ripe Grape Berry Lambda Triplex2 Library Vitis vinifera cDNA clone B3CS57RB007E11 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Lazge-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                              Vitis vinifera
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CN546489.1 GI:46911114
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genomics and Genetics Institute, GreenGene Biotech Inc., of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Nahm B.H
                                                                                   Universite de Bordeaux I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (bases 1 to 21)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gin, Kyeonggi, Koi
: 82 31 330 6193
: 82 31 321 6355
                                       Avenue Edouard Bourleaux,
00-33-(0)5-57-12-25-50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: pCR4-TOPO; Site 1: EcoRI; Callus was treated with ABA(20um) for 1hr. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was derived from rice Histone Deacetylase overexpression line."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /dev stage="proliferated callus on 2N6 media for 2 weeks" /lab_host="E.coli DH10B" /clone lib="OsHDAC1-overexpressing transgenic rice plasmid cDNA library (HD)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /db_xref="taxon:39947"
/clone="HD--08-C11"
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/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tissue_type="callus"
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Pred. No. 4
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                                                                                     Institut National de la Recherche
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21 bp
                                           BP 81,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          46;
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                                           33883 Villenave D'Ornon Cedex,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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VERSION
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AZ610868/c
LOCUS
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                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0436 row: G column: 1:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               plasmid inserts
Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                 Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    University of Utah Genome Center University of Utah
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igh quality sequence stop: 21.
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USA
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                                                        /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: pWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /dev stage="ripe stage"
/clome_lib="Ripe Grape Berry Lambda Triplex2 Library"
/note="Organ: Fruit without seeds; Vector: Lambda
TriplEx2; Site_1: SfiIA; Site_2: SfiIB; Oriented library"
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/mol type="mRNA"
/cultivar="Cabernet Sauvignon"
/db xref="Caxon:29760"
/clone="B3CS57RB007E11"
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                                           /mol_type="genomic DNA"
/strain="C578L/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                            organism="Mus musculus"
                                                                                                                                                                        sex="Male"
                                                                                                                                                                                           clone="UUGC1M0436G12"
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Pred. No. 46;
0; Mismatches
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RESULT 57
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Nouse, whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Ingert Length: 10000 Std Error: 0.00
Plate: 0560 row: D column: 04
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 801 585 5606
Fax: 801 585 7177
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University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
/lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/clone lib-"Mouse 10kb plasmid UUGCIM library"
/note="Vector: pMD42nv, Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
/http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
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                                                                                                                                                                                                                                                                                                           /mol_type="genomic DNA"
/strain="C578L/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                    clone="UUGC1M0560D04"
                                                                                                                                                                                                                                                                                                                                                                                              organism="Mus musculus"
                                                                                                                                                                                                                                                           sex="Male"
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Pred. No. 46;
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COMMENT

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BX556059/c
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1.1%; Score 15.8;
1 Similarity 89.5%; Pred. No. 48
17; Conservative
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BX556059 22 bp mRNA linear BST 10-OCT-200 BX556059 Glossina morsitans morsitans adult infected gut Glossina morsitans morsitans cDNA clone Tse24f09_plc, mRNA sequence.

BX556059
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             University of Wales,
Bangor LL57 2UW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Prof. M.J.Lehane
School of Biological Sciences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pathogen Sequencing Unit
The Sanger Institute The Wellcome Trust Genome Campus
Hinxton, Cambridge, CB10 1SA, UK
Request for clones, please contact: Mike Lehane
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Genome Biol. 4
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                                                                                                                                                                                                                                                                                                                                                                                                                       clones with suffix qlc are reverse primer reads starting of the cDNA all plc reads are from 3' end.
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                                                                                                       T.brucei"
                                                                                                                                                                          /clone="Tse24f09_p1c"
/tissue_type="adult infected gut"
/clone_lib="Glossina morsitans morsitans adult infected
                                                                                                                                                                                                                                                                                  /organism="Glossina morsitans morsitans"
/mol_type="mRNA"
/sub_species="morsitans"
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ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                               db_xref="taxon:37546"
                                                                                                                                                                                                                                                                                                                                                                                         ocation/Qualifiers
                                                                                                                               note="country: Zimbabwe; EST from adult gut infected with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 (10), R63 (2003)
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89.5%;
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  0; Mismatches
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Pred. No. 46;
                           48;
                                                DB 1; Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
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VERSION KEYWORDS SOURCE

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Genetics,

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CO778290
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CF310806/c
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                                                      Ambystoma mexicanum
Ambystoma mexicanum
Eukaryota; Metazoa; Chordata;
Eukaryota; Matrachia; Caudata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 22)
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oryza sativa (japonica cultivar-group) Oryza sativa (japonica cultivar-group) Oryza sativa (japonica cultivar-group) Eukaryota, Viridiplantae, Streptophyta, Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                             22 bp mRNA linear EST 05-AUG-2004
BL003B H01 6-Day Axolotl Tail Blastema (6DAxBL) Ambystoma mexicanum
CDNA 57 similar to hypothetical protein. mPNA accordance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CF310806 22 bp mRNA linear EST 15-AUG-2003 ABF--05-K20.gl ABF3-overexpressing transgenic rice plasmid cDNA library (ABF) Oryza sativa (japonica cultivar-group) cDNA clone ABF--05-K20, mRNA sequence.
                                                                                                                                                      EST
                                                                                                                                                                      cDNA 5' similar to hypo
CO778290
CO778290.1 GI:50994270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
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CF310806.1 GI:33682567
Habermann, B.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Nahm B.H.
                                           Атьувсота.
                                                                                                                             Ambystoma mexicanum (axolotl)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
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                    (bases 1 to 22)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note="Vector: pCR4-TOPO; Site 1: EcoRI; Leaf was dried for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /tissue_type="leaf"
/dev_stage="14 days after germination"
/lab_host="E.col1 DH10B"
/clone lib="ABF3-overexpressing transgenic rice plasmid cDNA library (ABF)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            db_xref="taxon:39947"
clone="ABF--05-K20"
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Bebin, A.G.,
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Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           bhnahm@bio.myongji.ac.kr
Herklotz, S.,
                                                             Craniata; Vertebrata; Euteleostomi; Salamandroidea; Ambystomatidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        48;
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Volkmer, M., Eckelt, K.,
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AZ823875
                                                                                                                                                                                                                                                                                        COMMENT
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               Tel: 801 585 5606
FAX: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0
Plate: 0098 row: K column: 07
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 22.
                                                                                                                                                                                                                                                                                                                                                                        Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 22)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AZ823875 22 bp DNA linear GSS 20-FEB-200 2M0098K07F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0098K07 F, genomic survey sequence.
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                                                                                                                                                                                                                    University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Tel: 0049 351 210 2620
Fax: 0049 351 210 1489
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                                                                                                                                                                                                                                                                                                             plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                        Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus (house mouse)
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Seq primer: GCA CAT TAG GCC TAT
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                                                                                                                                                                                                    84112,
                                                                                                                                                                                                                                                                                          Contact: Robert B. Weiss
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/cell type="regenerating tail blastema"
/cell type="regenerating tail blastema"
/clone_lib="6-Day Axolot! Tail Blastema (6DAxBL)"
/note="Vector: pCMVSport6; Site=1: Not1; Site=2: Sall;
/normalized cDNA plasmid library prepared by Invitrogen.
Vinormalized cDNA plasmid library prepared by Invitrogen.
Size fractionated mRNA was polydT primed and cloned into
Not1-Sall site of pCMVSport6. Bacterial host is
EMDH10B-TONA. Average insert size is 1.67 kB.
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/mol_type="mRNA"
/db_xref="taxon:8296"
    Location/Qualifiers
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Pred. No. 48;
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AUTHORS
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BH000233
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Best Local Similarity
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                                                                                     Tel: 801 585 5606
Feax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0287 row: L column: 21
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                             Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                         plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Verrebrata; Eureleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 22)
1 (bases 1 to 22)
1 (bases 1, base), Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BH000233 22 bp DNA linear GSS 27-APR-200200287L21R Mouse 10kb plasmid UUGC2M library Mus musculus genomic clone UUGC2M0287L21 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4 AAAAAAAAAAAAAAAAAG 22
                        High quality sequence stop: 22.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                     Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BH000233.1 GI:13871459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BH000233
                                                                    Class: plasmid ends
                                                                                                                                                                                                                                                              308,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi | 4732114 | gb | AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                           USA
                                                                                                                                                                                                                                                            Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA f
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         aboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  clone="UUGC2M0098K07"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15.8; DB 1; Pred. No. 48;
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AUTHORS
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ORGANISM
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VERSION
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AI708831/c
LOCUS
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                                                                         FEATURES
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Best Local
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1520 AAAAAAAAAAAAGTAAAAG 1538
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                                                                                                                                                                             Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Possible reversed clone: similarity on wrong strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 22)

Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie,T., Waterston,R. and Wilson,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AI708831 22 bp mRNA linear EST 04-JUN-1999 as27d12.x1 Barstead aorta HPLRB6 Homo sapiens cDNA clone IMAGE.2318423 3' similar to TR:Q33563 Q35563 EATRO 164 KINETOPLAST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 AAAAAAAAAAAAAAAAA 21
                                                                                                                         Seg primer: -40UP from Gibco
High quality sequence store
                                                                                                                                                                                                                                                                                                                                                                                                            Washington University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unpublished (1997)
Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WashU-NCI human EST Project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens (human)
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                                                                                           quality sequence stop: 1.
Location/Qualifiers
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314 286 1810
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMs92 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone_lib="Mouse 10kb plasmid_UUGC2M_library"
/note="Vector: PWD42nv; Purified genomic_DNA_from M.
musculus_C57BL/6J (female) was obtained from the Jackson
Laboratory Mouse_DNA_Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
organism="Homo sapiens"
on__type="mRNA"
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/strain="C57BL/6J"
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Pred. No. 48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dubuque, T., Geisel, G., Jost, S.,
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JOURNAL COMMENT
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SOURCE
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AJ747407
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VERSION
     DEFINITION
                                                   AW332181/c
                                                                              RESULT 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE
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Best Local
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Best Local Similarity
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AW332181 22 bp mRNA linear EST S5C7 AGS-1 Pneumocystis carinii cDNA 3', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AJ747407
AJ747407 forward - stimulated minus unstimulated macrophage scrofa cDNA clone F_C0001825b_A06, mRNA sequence.
AJ747407
AJ747407
AJ747407.1 GI:49917653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Hopwood PA

Contact: Hopwood PA

Dept. of Preclinical Veterinary Sciences

Royal School for Veterinary Studies

Summerhall, Edinburgh, EH9 1QH, UNITED KINGDOM

Sequencing was performed by ARK genomics. This clone is available

from ARK- Genomics, Roslin Institute, Roslin, Midlothian EH25 9PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Development of a porcine cDNA microarray Unpublished (2004)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bukaryota; Metazoa; Chordata; Craniata; Vertabrata; Buteleostomi;
Mammalia; Butheria; Cetartiodactyla; Suina; Suidae; Sus.
1 (bases 1 to 22)
Hopwood, P.A., Zhang, F., Lowden, S., Talbot, R., Burt, D., Archibald, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sus scrofa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and Dixon,L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sus scrofa (pig)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TTTTTGTGGTGCGTGCGTGC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       See www.ark- genomics.org or contact info@arkgenomics.org.
Location/Qualifiers
                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="mRNA"
/db xref="taxon:9823"
/db xref="taxon:9823"
/clone="F_C0001825b_A06"
/tissue_type="lung"
/cell_type="macrophage"
/clone_lib="forward - stimulated minus unstimulated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      organism="Sus scrofa"
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/lab_host="DH10B (phage resistant)"
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                                                                                                                                                                                                                                                                                       1.1%;
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                                                                                                                                                                                                                                                                                       Score 15.6;
Pred. No. 54
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Pred. No. 54;
                                                                                                                                                                                                                                                                    Mismatches
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                             EST 31-JAN-2000
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S8A2 AGS-1 Pneumocystis carinii
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University of Kentucky
101 Morgan Building, University of Kentucky, Lexington,
40506-0225, USA
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Eukaryota; Fungi; Ascomycota; Pneumocystidomycetes;
Pneumocystidaceae; Pneumocystis.
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Smullan,A.G., Arnold,J., Weise,M., Wunderlich,J.,
Edman,J.C., Kovacs,J. and Cushion,M.
Expressed sequence tags from Pneumocystis carinii
Unpublished (2000)
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University of Kentucky
101 Morgan Building, Universi
40506-0225, USA
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Smulian,A.G., Arnold,J., Weise,M., Wunderlich,J., Edman,J.C., Kovacs,J. and Cushion,M.
Expressed sequence tags from Pneumocystis carinii
                                                                                                                                                                                Tel: 606 257 2161
Fax: 606 257 1717
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Contact: Staben C
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AW332181.1
                                                                                                                               Email: staben@pop.uky.edu.
Location/Qualifiers
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Location/Qualifiers
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/organism="Pneumocystis
/mol_type="mRNA"
/db xref="taxon:4754"
/lab_host="E. coli"
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/db_xref="taxon:4754"
/lab_host="E. coli"
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oryza sativa (japonica cultivar-group)
oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
CF300133 22 bp mRNA linear EST 15-AUG-2003
7LEAF--04-G19.bl Rice leaf plasmid cDNA library II (7LEAF) Oryza
8ativa (japonica cultivar-group) cDNA clone 7LEAF--04-G19, mRNA
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Le Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.

Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)

Contact: Nahm B.H.
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7LEAF--03-F06.bl Rice leaf plasmid cDNA library II (7LEAF) Oryza
8ativa (japonica cultivar-group) cDNA clone 7LEAF--03-F06, mRNA
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CF299342.1 GI:33671103
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jin, Kyeonggi, Korea
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/note="Vector: Lambda ZAP II; Site_1: EcoRI; Site_2: XhoI;
/note="Vector: Lambda ZAP II; Site_1 = EcoRI; Site_2: XhoI;
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/note="Vector: Lambda ZAP II; Site_1: EcoRI; Site_2: ABC II;
/note="Vector: Lambda II; Site_1: EcoRI; Site_2: ABC II; Site_1: ABC II; Sit
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/dev_stage="7 days after germination"
/lab_host="E.coli DH108"
/lab_host="E.coli DH108"
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/clone ib="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped
/mche="Vector: pCR4-TOPO; Site_1 seconi; mRNA was capped
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/mol_type="mRNA"
/cultivar="Nackdong"
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'clone="7LEAF--03-F06"
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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ABF--04-P14.gl ABF3-overexpressing transgenic rice plasmid cDNA
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., L
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                              Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                   Genomics and Genetics Institute, GreenGene Biotech Inc.; of Bioscience and Bioinformatics, MyongJi University
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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RT-PCR."
/organism="Oryza sativa
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/mol_type="mRNA"
/cultivar="Nackdong"
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Pred. No. 54;
                                                                         bhnahm@bio.myongji.ac.kr
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  (japonica cultivar-group)"
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Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.

1 (bases 1 to 22)
1 (bases 1 to 22)
1 (bases 1 to 21)
1 (Song,S.I., Kim,J.K., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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22 bp mRNA linear EST 15-AUG-2003
ABF--06-G21.gl ABF3-overexpressing transgenic rice plasmid cDNA
library (ABF) Oryza sativa (japonica cultivar-group) cDNA clone
ABF--06-G21, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea Tel: 82 31 330 6193 Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Nahm B.H.
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Location/Qualifiers
                                                                                         /note="Vector: pCR4-TOPO; Site 1: EcoRI; Leaf was dried for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression line."
                                                                                                                                                                                                        /tissue_type="leaf"
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/lab_host="E.coli DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /tissue_type="leaf"
/dev stage="14 days after germination"
/lab_host="E.coli DH10B"
/clone lib="ABF3-overexpressing transgenic rice plasmid
/clone library (ABF)"
                                                                                                                                                                                                                                                                                                                                                                       /db_xref="taxon:39947"
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/clone="ABF--04-P14"
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'cultivar="Nackdong"
                     1.1%;
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                    Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
1 (bases 1 to 22)
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptphyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Lazge-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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CF312498.1 GI:33684259
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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CF311713.1 GI:33683474
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Location/Qualifiers
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ABF--07-B13, mRNA sequence.
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Similarity 81.8%;
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  J.S., Jun, K.M., Cheong, P.J.,
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/dev stage="14 days after germination"
/lab_host="E.coli DH10B"
/clone lib="ABF3-overexpressing transgenic rice plasmid cDNA llbrary (ABF)"
/note="Vector: pCR4-TOPO; Site 1: EcoRI; Leaf was dried for 2hrs. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was prepared from ABA-responsive element binding transcription factor 3 overexpression
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/clone="ABF--07-B13"
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/cultivar="Nackdong"
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Pred. No. 54;
  Kim, M.J.,
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Lee, T.H., Shin, Y.C.,
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CF330679/c
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
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NACL--06-H22.bl Rice callus plasmid cDNA library (NACL) Oryza
Bativa (japonica cultivar-group) cDNA clone NACL--06-H22, mRNA
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                                                                                                                                                                                                                           Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea Tel: 82 31 330 6193 Fax: 82 31 321 6355
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
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                                                                                                                                                                                                                                                                                                                                   Contact: Nahm B.H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Nahm B.H.
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82 31 321 6355
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Location/Qualifiers
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/lab_host="B.coli DH10B"
/clone_lib="ABF3-overexpressing transgenic rice plasmid
cDNA library (ABF)"
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|mol type="mRNA"
|cultivar="Nackdong"
|db xref="taxon:39947"
|clone="ABF--08-E15"
                                  /db_xref="taxon:39947"
/clone="NACL--06-H22"
/tissue_type="callus"
dev_stage="proliferated/
lab_host="E.coli DH10B"
                                                                                               organism="Oryza sat
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Pred. No. 54;
0; Mismatches
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CF334781 22 bp mRNA linear EST 18-AUG-20 JMT--04-D05.gl AtJMT-overexpressing transgenic rice plasmid cDNA library (JMT) Oryza sativa (japonica cultivar-group) cDNA clone JMT--04-D05, mRNA sequence.
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Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,

Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.

Large-scale Sequencing Analysis of Rice ESTs

Unpublished (2003)
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Oryza sativa (japonica cultivar-group)
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
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/dev stage="14 days after germination"
/lab_host="E.coli DH10B"
/clone lib="AtDWT-overexpressing transgenic rice plasmid cDNA library (JMT)"
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/note="Vector: pCR4-TOPO; Site 1: EcoRI; mRNA was capped
with oligoribonuclectides and then used as templates for
RT-PCR."
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/mol_type="mRNA"
/cultivar="Nackdong"
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/clone="JMT--02-F04"
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Best Local Similarity
Matches 18; Conser
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source
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantea; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.

1 (bases 1 to 22)
Kim, J. S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Shin, Y.C.,
Song, S. I., Kim, J. K., Kim, Y.-K. and Nahm, B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                  Oryza sativa (japonica cultivar-group) oryza sativa (japonica cultivar-group) oryza sativa (japonica cultivar-group) Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Ehrhartoideae; Oryzeae; Oryza.
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                   Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@gbio.com, bhnahm@bio.myongji.ac.kr.
                                                                                                                                                                                                              1 (bases 1 to 22)
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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Location/Qualifiers
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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EST.
                                                                                                                                                                                             Contact: Nahm B.H.
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/dev stage="14 days after germination"
/lab_host=="E.coli DH10B"
/clone lib="AtJMT-overexpressing transgenic rice plasmid cDNA library (JMT)"
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/mol_type="mRNA"
/cultivar="Nackdong"
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clone="JMT--04-D05"
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Pred. No. 54;
O; Mismatches
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transgenic rice plasmid cDNA
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RESULT 77
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                  Query Match
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Nahm B.H.
Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 22)
Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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Location/Qualifiers
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/clone lib-"AtcMT-overexpressing transgenic
cDNA library (JMT)"
                                                                                            CDNA library (JMT)"

/note="Vector: pCR4-TOPO; Site_1: EcoRI; Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was pepared from Arabidopsis Jasmoinate Carboxyl
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pepared from Arabidopsis Jasmoinate Carboxyl
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/cultivar="Nackdong"
/db_xref="taxon:39947"
                                                                            methyltransferase overexpression line."
                                                                                                                                                                        /dev_stage="14 days after germination"
/lab_host="E.coli DH10B"
/clone_lib="AtJMT-overexpressing transgenic rice plasmid
                                                                                                                                                                                                                                                                /mol_type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:39947"
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                                                                                                                                                                                                                                                      clone="JMT--08-B11"
                1.1%;
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                                    Score 15.6;
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AUTHORS
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Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
1 (bases 1 to 22)
Abball P., Agasse A., Ageorges, A., Atanassova, R.,
Couture C., Dedaldechamp, F., Delrot, S., Glissant,
Hamdi, S., Romieu, C. and Terrier, N.
                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; Vitaceae; Vitis.
                                                                                                                                                                                                           Vitis vinifera
Vitis vinifera
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Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae,
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RCL1--01-P07.gl Regenerated callus lambda phage cDNA library (RCL1)
Oryza sativa (japonica cultivar-group) cDNA clone RCL1--01-P07,
                                                                                                                                                                                                                                                                                                                                                                                                                             CN545550
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Nahm B.H
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/mol_type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:39947"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'Clone_lib="Regenerated callus lambda phage cDNA library
|RCL1)|
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      dev_stage="proliferated callus on 2N6 media for 30 days"
|lab_host="E.coli SOLR"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tissue_type="callus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    clone="RCL1--01-P07"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             81.8%;
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                                                                                                                                                                                                                                                                                                                                                              Skin
3', m
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15.6;
Pred. No. 54
                                                                                                                                                                                                                                                                                                                                                           mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         bhnahm@bio.myongji.ac
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                            Barrieu, F.,
D., Grimple
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                               Grimplet, J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
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VERSION
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AZ310066/c
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AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
                                                                                                                                                                                                                                                                                                                                                              Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AZ310066.1 GI:10351682
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Agronomique
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Similarity 81.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                      Location/Qualifiers
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   db_xref="taxon:10090"
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TITLE
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
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Contact: Hamdi S.
UMR 619 - Equipe Biologie de la Vigne
Universite de Bordeaux I, Institut National de la Recherche
                                                                                                                                                                                                                                                                                                                                                            Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AZ310066 22 bp DNA linear GSS 2: 1M0018D18R Mouse 10kb plasmid UUGCIM library Mus musculus clone UUGC1M0018D18 R, genomic survey sequence.
                                                                                                                                                                                                     Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0018 row: D column: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                    Seq primer: CACACAGGAAACAGCTATGACCClass: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: s.hamdi@bordeaux.inra.fr
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                                                                                                                              quality sequence stop: 22.
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/clome_lib="Ripe Grape Skin Triplex2 Library"
/note="Organ: Fruit skin; Vector: Lambda TriplEx2; Site_1:
SfiIA; Site_2: SfiIB; Oriented library"
/mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="mRNA"
/culfivar="Cabernet Sauvignon"
/db xref="raxon:29760"
/clone="B3CS00RL003D05"
                                                   organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            organism="Vitis vinifera"
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                                                                                                                                                                                                                                Std Error: 0.00
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Query Match
Best Local Similarity
Matches 18; Conserv
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                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0034 row: I column: 16
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        plasmid inserts
Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 22)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AZ316361 22 bp DNA linear GSS 29-SEP-2001M0034I16F Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGC1M0034I16 F, genomic survey sequence.
                                                                                                                      High quality sequence stop: 22.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AZ316361.1 GI:10364110
                                                                                                                                                                              Class: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clome_lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                clone="UUGC1M0018D18"
                  mol_type="genomic DNA"
strain="C57BL/6J"
                                                                     organism="Mus musculus"
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xref="taxon:10090"

 Mismatches

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Pred. No. 54;
                                                                                                                                                                                                                                                      Std Error: 0.00
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ORGANISM
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AZ317017/c
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RESULT 81 AZ316361

SOURCE ORGANISM

KEYWORDS VERSION ACCESSION DEFINITION

REFERENCE

AUTHORS

COMMENT

JOURNAL

FEATURES

/db_xref="taxon:10090" /clone="UUGC1M0035P09"

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1516 AATTAAAAAAAAAAAAAGTAAAA 1537
                                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error: 0.
Plate: 0035 row: P column: 09
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 22.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AZ317017 22 bp DNA linear GSS 29-SEP-200 1M0035P09F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0035P09 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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/mol_type="genomic DNA'
/strain="C57BL/6J"
                                                                                                                      Location/Qualifiers
                                                        organism="Mus musculus"
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                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0089 row: E column: 07
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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AZ351527
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Fax: 801 585 7177
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Contact: Robert B. Weiss
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/note="Vector: PWD42nv; Purified genomic DNA f
                                                                                                     /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /lab_host="E. Coli strain XL10-Gold, T1-resistant,
                                                clone="UUGC1M0089E07"
                                                                                 db_xref="taxon:10090"
                                                                                                                                                                 organism="Mus musculus"
host="E. Coli strain XL10-Gold, T1-resistant,
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Insert Length: 10000 Std Error: 0
Plate: 0099 row: M column: 15
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 22.
Location/Qualifiers
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Fax: 801 585 7177
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Contact: Robert B.
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/note="Vector: PWD42nv; Purified genomic DNA f
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'lab_host="E. Coli strain XL10-Gold, T1-resistant,
'clone_lib="Mouse 10kb plasmid UUGCIM library"
                                                                                            sex="Male"
                                                                                                                                                  clone="UUGC1M0099M15"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.6;
Pred. No. 54;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2030 E., SLC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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ACCESSION VERSION

KEYWORDS

RESULT 84 AZ357630 LOCUS

DEFINITION

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Matches

REFERENCE AUTHORS

TITLE

JOURNAL

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JOURNAL COMMENT
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AUTHORS
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ORGANISM
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VERSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DEFINITION
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Best Local Similarity
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1M0147N14R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0147N14 R, genomic survey sequence.
AZ388103
                                                                                                                                                                                                                                                                           High quality sequence stop: 22.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0147 row: N column: 14
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (Dases 1 to 22)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plasmid inserts
Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          308,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Laboratory Mouse DNA Resource
/lab_host="E. Coli strain XL10-Gold, T1-resistant, F
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                               mol_type="genomic DNA"
strain="C57BL/6J"
                                                                                                                                                                                                                        organism="Mus musculus"
                                                                                        sex="Male"
                                                                                                               clone="UUGC1M0147N14"
                                                                                                                                           db_xref="taxon:10090"
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Pred. No. 5
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KEYWORDS

RESULT 85 AZ388103

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Matches

RESULT 86

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SOURCE
ORGANISM
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LOCUS
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Best Local 8
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                                                                                                                                                                                                                                                                                                                                            Tel: 801 585 5606
FAX: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.0
Plate: 0168 row: P column: 24
Seg primer: CACAGGAAACAGCTATGACC
Class: plasmid ends
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1M0168P24R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0168P24 R, genomic survey sequence
AZ401908
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                  High quality sequence stop: 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               308,
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(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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/lab_host="E. Coli strain XL10-Gold, T1-resistant,
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from N
                                                                                                                                          /mol_type="genomic_DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                        sex="Male"
                                                                                                               clone="UUGC1M0168P24"
                                                                                                                                                                                                                                  organism="Mus musculus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15.6;
Pred. No. 54;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2030 E., SLC,
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KEYWORDS VERSION ACCESSION

JOURNAL COMMENT

TITLE

FEATURES

C57BL/6J (male) was obtained

REFERENCE AUTHORS

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REFERENCE
AUTHORS
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Wahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0203 row: A column: 24
Seq primer: CACACAGGAAAACAGCTATGACC
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1M0203A24R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0203A24 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
Contact: Robert B. Weiss
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                                                                                                                                                                                                                                                                                                                                                                               ass: plasmid ends
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
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/lab_host="B. Coli strain XLI0-Gold, Tl-resistant, /Clone_lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: pW042nv; Purified genomic DNA from hmusculus C57BL/6J (male) was obtained from the Jac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Laboratory Mouse DNA Resource
                                                                                                                                                                                               mol_type="genomic DNA"
'strain="C57BL/6J"
                                                                                                                    sex="Male"
                                                                                                                                                  'db_xref="taxon:10090"
'clone="UUGC1M0203A24"
                                                                                                                                                                                                                                                              organism="Mus musculus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. Mouse whole genome scaffolding with paired end reads from 10kb Mouse whole genome scaffolding with paired end reads
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0212 row: A column: 05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 22)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                       Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        University of Utah
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plasmid inserts
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SA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of phasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone_lib="Mouse_10kb_plasmid_UUGC1M_library" /note="Vector: PwD42nv; Purified_genomic_DNA_from M. musculus_CS7BL/6J (male) was obtained_from the Jackson_Laboratory_Mouse_DNA_Resource_thtp://www.jax.org/resources/documents/dnares/). The DNA_from the Laboratory_mouse_DNA_resources/documents/dnares/).
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                                                                                                                                                                                                                                      /mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                                                                                    /db_xref="taxon:10090"
/clone="UUGC1M0212A05"
                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                          organism="Mus musculus"
                                                                                                                                                              sex="Male"
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Mouse, whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0264 row: G column: 12
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (2000)
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AZ459654.1 GI:10617779
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert B. Weiss
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/lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clone lib="Mouse 10kb plasmid UUGC1M library"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/63 (male) was obtained from the Jacksor Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The I
                                                                                                                                                                                       /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0264G12"
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Pred. No. 54;
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                                                              from the Jackson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0272 row: E column: 24
Seg primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030
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1M0272E24F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
Clone UUGC1M0272E24 F, genomic survey sequence.
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Unpublished (2000)
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High quality sequence stop: 22.
Location/Qualifiers
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                             /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lbb="Mouse 10kb_plasmid_UUGCIM_library"/note="Vector: pwp82ny; Purified genomic_DNA from M.musculus C57BL/6J (male) was obtained from the Jackso)
Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                                                                                           /db_xref="taxon:10090"
/clone="UUGC1M0272E24"
                                                                                                                                                                                                                                                                                                                   mol_type="genomic DNA"
strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                        organism="Mus musculus"
                                                                                                                                                                                                                                   sex="Male"
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31.8%; Pred. No. 54;
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0; Mismatches
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Best Local Similarity
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Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0272 row: E column: 12
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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1M0272E12R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
Clone UUGC1M0272E12 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           plasmid inserts
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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/lab host-"E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib-"Mouse 10kb plasmid UUGCIM library"
/note-"Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
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/clone="UUGC1M0272E12"
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/strain="C57BL/6J"
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Pred. No. 54;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
Plate: 0374 row: J column: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mouse whole genome scaffolding with paired end reads from 10kb
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant vellocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Biomedical Polymers Research Bldg., 20 S.
                                                                                                                                                                                                                                                              /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /clone_lib="Mouse_10kb_plasmid_UUGC1M_library" /note="Vector: PW042Tv; Purified_genomic_DNA_from_M.musculus_C57BL/6J_(male) was obtained_from_the_Jackson_musculus_C57BL/6J_(male) was obtained_from_the_Jackson_Musculus_C57BL/6J_(mal
                                                                                                                                                                                                                          Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /db_xref="taxon:10090"
/clone="UUGC1M0374J15"
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                                                                                                                                                                                                                                                                                                                                                                                                                                               /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /strain="C57BL/6J"
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Pred. No. 54;
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0430 row: A column: 13
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Fax: 801 585 7177
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/lab host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone lib="Mouse 10kb plasmid UUGCIM library"
/clone="Vector: PMD42nv; Purified genomic DNA from M.
musculus C57BL/6y (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
                                                                                                                                                                                                                                                                                                                                                                            /mol_type="genomic DNA"
/strain="C57BL/6J"
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Pred. No. 54
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VERSION KEYWORDS RESULT 94 AZ654691 FEATURES REFERENCE ACCESSION DEFINITION JOURNAL TITLE ORGANISM Matches Query Match Best Local Similarity source 1516 AATTAAAAAAAAAAAAGTAAAA 1537 18; 1 AAAAAAAAAAAAAAAAAAAAA 22 Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Err Plate: 0529 row: D column: 0 Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Welss, R.
Mouse whole genome scaffolding with paired end reads from 10kb Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus. 1 (bases 1 to 22) AZ654691 22 bp DNA linear GSS 14
1M0529D05F Mouse 10kb plasmid UUGC1M library Mus musculus
clone UUGC1M0529D05 F, genomic survey sequence. Class: plasmid ends
High quality sequence stop: 22.
Location/Qualifiers Seq primer: CGTTGTAAAACGACGGCCAGT University of Utah Genome Center University of Utah Unpublished (2000) Dunn, D., Aoyagi, A., Barber, M., Tel: 801 585 5606 Fax: 801 585 7177 Contact: Robert B. Weiss plasmid inserts Mus musculus Mus musculus (house mouse) AZ654691.1 AZ654691 308, USA Conservative 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLL10-Gold (Stratagene) cells and selected for ampicillin resistance." (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource Biomedical Polymers Research Bldg., 20 /organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J" adaptored DNA was purified and size-selected for a 9.5 to db_xref="taxon:10090" sex="Male" clone="UUGC1M0529D05" GI:11791837 1.1%; Score 15.6; 81.8%; Pred. No. 54; 0; Mismatches Std Error: Beacorn, T., Duval, B., Hamil, C., 0.00 DB 1; 4. Length 22; s. 2030 E., SLC, GSS 14-DEC-2000 0 Gaps genomic S 0

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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Nouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error:
Plate: 0554 row: A column: 24
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
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GSS.
Mus musculus (house mouse)
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1M0554A24F Mouse 10kb plasmid UUGCIM library Mus musculus genomic
Clone UUGC1M0554A24 F, genomic survey sequence.
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Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA (http://www.jax.org/resources/documents/dnares/). The shared by a hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel
                                                                                                                                                                                                                                                              /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" (clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: pW042nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
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                                                                                                                                                                                                                                                                                                                                                                                                       /db_xref="taxon:10090"
/clone="UUGC1M0554A24"
                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    organism="Mus musculus"
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                                                                                                                                                                                                                                                                                                                                                                                sex="Male"
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; Pred. No. 54;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0016 row: I column: 12
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Unpublished (2000)
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801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative
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                                                                                                                                                                                                                                                                                    /clone_lib="Mouse_10kb_plasmid_UUGC1M_library"
/note="Vector: PWD42nv; Purified_genomic_DNA_from_M.
musculus_C57BL/6J_(male)_was_obtained_from_the_Jackson
                                                                                                                                                                                                                                                             Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mol_type="genomic DNA"
/strain="C57BL/6J"
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81.8%;
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Mouse, whole genome scaffolding with paired end reads from 10kb
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Insert Length: 10000 Std Error: C
Plate: 0028 row: E column: 04
Seq primer: CACACAGGAAACAGCTATGACC
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1 (bases 1 to 22)
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2M0028E04R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0028E04 R, genomic survey sequence.
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Fax: 801 585 7177
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Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA (http://www.jax.org/resources/documents/dnares/). The Shange through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gi|4732114|gb|AF129072.1), a copy-number
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                                                                                                                                                                                                                                                                                           /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mol_type="genomic_DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                       clone="UUGC2M0028E04"
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Pred. No. 54;
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inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

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REFERENCE AUTHORS VERSION KEYWORDS FEATURES COMMENT SOURCE DEFINITION AZ787098/c LOCUS RESULT ACCESSION TITLE ORGANISM Matches Query Match Best Local (JOURNAL source 1516 AATTAAAAAAAAAAAAGTAAAA 1537 22 AAAAAAAAAAAAAAAAAAAA 1 22 bp DNA linear GSS 1t 2M0033A05F Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC2M0033A05 F, genomic survey semience AZ787098 Seq primer: CGTTGTAAAACGACGGCCAGT Class: plasmid ends High quality sequence stop: 22. University of Utah Genome Center University of Utah Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rokes, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 22) 18; Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Err Plate: 0033 row: A column: 0 plasmid inserts Mus musculus Tel: 801 585 5606 Fax: 801 585 7177 Contact: Robert Unpublished (2000) Mus musculus (house mouse) AZ787098.1 Similarity 308, Conservative USA (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /Clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Biomedical Polymers Research Bldg., 20 Laboratory Mouse DNA Resource /db_xref="taxon:10090" /clone="UUGC2M0033A05" /mol_type="genomic DNA" /strain="C57BL/6J" Location/Qualifiers organism="Mus musculus" 1.1%; Score 15.6; 81.8%; Pred. No. 54; Weiss 0 Std Error: 0.00 Mismatches DB 1; 4. Length 22; <u>ა</u> 2030 GSS 16-FEB-2001 SLC, S 0

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AZ787606/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0034 row: G column: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                University of Utah Genome Center University of Utah Rm. 308 P4-----
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2M0034G12F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
Clone UUGC2M0034G12 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
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Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Conservative
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pUM42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   organism="Mus musculus"
/mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            gex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    clone="UUGC2M0034G12"
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Pred. No. 54;
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Insert Length: 10000 Std Err
Plate: 0045 row: A column: 2
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Fax: 801 585 7177
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University of Utah
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWH92 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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                                                                                                                                                                                                                                                                                                                                                    /lab host="B. Coli strain XL10-Gold, T1-resistant, F-"/Clome_lib="Mouse 10kb plasmid UUGCIM library"/note="Vector: pwp82nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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                                                                                                                                                                                                                                                                                                                                  Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mol_type="genomic
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               clone="UUGC2M0045A24"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         db_xref="taxon:10090"
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Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.6; DB 1;
Pred. No. 54;
0; Mismatches 4;
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                                                                                                                                                                                                                                                                                                                                                         Jackson
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AZ810674/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AZ810674 22 bp DNA linear GSS 20 2M0076E19F Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC2M0076E19 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0076 row: E column: 19
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 22)
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GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           quality sequence stop: 22.
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              (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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adaptored vector DNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  clone="UUGC2M0076E19"
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/strain="C57BL/6J"
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  transformed into
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Query Match
                                          chemically-competent E. coli XL10-Gold and selected for ampicillin resistance.
Score 15.6; 1
               DB 1;
                                                  (Stratagene) cells
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RESULT 102
AZ820439/c
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Class: plasmid ends
High quality sequence stop: 22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0092 row: K column: 13
Seq primer: CACACAGGAAACAGCTATGACC
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse whole genome scaffolding with paired end reads from 10kb
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                / Tab host="E. Coli strain Xi10-Gold, T1-resistant, F-"
//lab host="E. Coli strain Xi10-Gold, T1-resistant, F-"
//clone lib="Mouse 10kb plasmid UUGCIM library"
//note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
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adaptored vector DNA, ar chemically-competent E.
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/clone="UUGC2M0092K13"
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/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sex="Male"
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AZ841661
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Location/Qualifiers
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Plate: 0139 row: I column: 18
Seg primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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Fax: 801 585 7177
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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ilarity 81.8%;
Conservative
                                                                       (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pumple (gi | 4732114 | gb | AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/strain="C57BL/6J"
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Nouse, Whole genome scaffolding with paired end reads from 10kb Mouse, whole genome scaffolding with paired end reads
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Location/Qualifiers
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Insert Length: 10000 Std Error:
Plate: 0142 row: I column: 24
Seq primer: CGTTGTAAAACGACGGCCAGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of phsmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA and transformed into
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/strain="C57BL/6J"
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Pred. No. 54;
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AZ946102
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AZ946102 22 bp DNA linear GSS 27-APR-200
2M0207D13R Mouse 10kb plasmid UUGC2M library Mus musculus genomic
clone UUGC2M0207D13 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0207 row: D column: 13
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus
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                                                                                                              adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored.
                                               adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                 (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant vellocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /lab host="E. coli strain XL10-Gold, T1-resistant, F-"
/clome_lb="Mouse 10kb plasmid UUGC2M library"
/note="Vector: PWPA2nv; Purified genomic DNA from M.
musculus C57BL/6J (female) was obtained from the Jack:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'sex="Female"
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/strain="C57BL/6J"
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Pred. No. 5;
  Score 15.6;
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Matches 18; Conserv
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                                                                       Trypanosoma brucei
Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               insert libraries for whole genome shotgun sequencing projects. In Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).

Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                to give a tight size distribution ( 4~\rm kb) . The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nhl@sanger.ac.uk
Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically shea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 15A, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
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Hall, N., Bowman, S., Lennard, N.J., Doggett, J., Atkin, R., Chillingworth, C., Ormond, D., Harris, B., El-Sayed, N., Hou, L.,
                                                                                                                                      GSS.
                                                                                                                                                                                                              T. brucei sheared genomic
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                                                                                                                   Trypanosoma brucei
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/strain="TREU927"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone="131b09"
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Pred. No. 54;
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A clone 131b09, forward sequence,
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                                                                                                                                                                                                                                                                                          Direct Submission
Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing
project Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,
Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
nhl@sanger.ac.uk
Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
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4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                 Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                          to give a tight size distribution (
4 kb). The v + i method used for the library construction is
described in detail in Smith, H. and Venter, J.C. (Making small
insert libraries for whole genome shotgun sequencing projects.
Genome Sequencing: A Practical Approach, eds. M. Vaudin and B.
Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
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Trypanosoma brucei
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AL454256.1 GI:11855060
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Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing
Submitted (10-DEC-2000) Trypanosoma brucei genome Campus, Hinxton,
Cambridge CB10 18A, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Trypanosoma.
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/mol_type="genomic DNA"
/strain="TREU927"
/organism="Trypanosoma brucei"
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/strain="TREU927"
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Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
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4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Bartell, Oxford University Press, 1999).
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                                                            AG194579
AG194579.1
                                                                                                                         Pan troglodytes DNA, clone: RP43-072N05.TJ,
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Pan troglodytes (chimpanzee)
Pan troglodytes
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/clone="35c12"
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/clone="380a07"
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/strain="TREU927"
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                                                                                                                                                                                                                                                                             Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplanta; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
1 (bases 1 to 19)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    CF334610 19 bp mRNA linear EST 18-AUG-2003 JMT--03-P13.bl AtJMT-overexpressing transgenic rice plasmid cDNA library (JMT) Oryza sativa (japonica cultivar-group) cDNA clone JMT--03-P13, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted (07-JAN-2002) Hong-Seog Park, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Genome Research Center (GRC); 52, Oun-dong, Yusong-gu, Daejeon 305-333, Korea (E-mail:redstone@mail.kribb.re.kr, URL:http://phs.grc.kribb.re.kr/, Tel:82-42-866-7181, Fax:82-42-860-4409) Clones are derived from the chimpanzee BAC library RP-43 This BAC end was generated during the R&D process and may have higher chance
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                                                    Email: bhnahm@ggbio.com, bhn.
Location/Qualifiers
                                                                                                                                                                                                     Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                CF334610 GI:33817556 CF334610.1 GI:33817556 EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Park,H., Kim,Y., Kim,S., Han,Y., Woo,T., Park,K., Eun,C.J., Hoon,S.T., Chu,M., Kim,H., Joo,S., Kim,C., Song,W. and Yoo,H. Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Park,H., Kim,Y., Kim,S., Han,Y., Woo,T., Park,K., Eun,C.J., Hoon,S.T., Chu,M., Kim,H., Joo,S., Kim,C., Song,W. and Yoo,H. BAC end sequences of Library RP-43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
                                                                                             Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                 Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                        Contact: Nahm B.H
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R.Site 1 : EcoRI
R.Site 2 : EcoRI.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               organism="Pan troglodytes"
/mol_type="genomic DNA"
/db_xref="taxon:9598"
/organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /cell_type="lymphocytes"
/clone_lib="RP-43 Chimpanzee Male BAC Library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.6;
Pred. No. 54;
                                                                         bhnahm@bio.myongji.ac.
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CL680297/c
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CF319122/c
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     CF319122
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1521 ААААААААААСТАААА 1537
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CL680297 CL680297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Evolutionary Biology
Max-Planck-Institute for Developmental Biology
Spemannstr. 37-39, Tuebingen D-72076, Germany
Tel: 00497071601371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pristionchus pacificus
Nucleic Acids Res. 32 (1), D421-D422 (2004)
Contact: Sommer RJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pristionchus pacificus
Eukaryota; Metazoa; Nematoda; Chromadorea; Diplogasterida;
Neodiplogasteridae; Pristionchus.
1 (bases 1 to 20)
                                                                                                                                                                                                                                                                                                                                                                                             Email: ralf.sommer@tuebingen.mpg.de
This library was generated at Caltech,
sequenced at Vancouver, Canada.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Srinivasan, J., Otto, G.W., AppaDB: an AcedB database
                                                                                                                                                                                                                                                                                                                                                          Class: fosmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fax: 00497071601498
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AAAAAAAAAAGAAAA
                                                                                Conservative
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/note="Vector: pcr4-TOPO; Site_1: EcoRI; Oligo-capped mRNA was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         was reverse transcribed and then used for PCR. pepared from Arabidopsis Jasmoinate Carboxyl methyltransferase overexpression line."
                                                                                                                                                                                                                                                /organism="Pristionchus
/mol_type="genomic DNA"
/strain="California"
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/lab_host="E.coli DH10B"
/clone_lib="AtJMT-overexpressing transgenic rice plasmid
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/db_xref="taxon:39947"
/clone="JMT--03-P13"
                                                                                                                                                              /note="Vector: pEpifos-5 Fosmid vector"
                                                                                                                                                                                   /clone_lib="Mixed stage
var. California"
                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                                                                                                                                                                                                                             db_xref="taxon:54126"
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                                                                                                   94.18;
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94.1%; Pred. No. 53;
ive 0; Mismatches
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                                                                                                 Score 15.4;
Pred. No. 56;
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for the nematode satellite
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                                                                                                                                                                                                       fosmid library of P.
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                                                                                                                     Length 20;
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21 bp

mRNA

linear

EST

15-AUG-2003

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T49097
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17
                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

(bases 1 to 25)

Hillier, L., Lennon, G., Becker, M., Bonaldo, M. F., Chiapelli, B., Chissoe, S., Dietrich, N., Dubuque, T., Favello, A., Gish, W., Chissoe, S., Dietrich, N., Kucaba, T., Lacy, M., Le, M., Le, N., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Parange, C., Rifkin, L., Rohlfing, T., Schallenberg, K., Soares, M. B., Tan, F., Thierry-Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P., Waterston, R., Wilson, R. and Marra, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oryza sativa (japonica cultivar-group)
oryza sativa (japonica cultivar-group)
oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                          749097
749097
755 bp mRNA linear EST 06-FEB-19
yb08h08.81 Stratagene placenta (#937225) Homo sapiens cDNA clone
IMAGE.70623 3' similar to similar to gb.X62744 CLASS II
HISTOCOMPATIBILITY ANTIGEN, M ALPHA CHAIN (HUMAN), mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
Email: bhnahm@ggbio.com, bhnahm@bio.myongji.ac.
Location/Qualifiers
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1 (bases 1 to 21)

Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le Kim,J.S., Jun,K.M., Kim,Y.-K. and Nahm,B.H. Song,S.I., Kim,J.K., Kim,Y.-K. and Rice ESTs
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HD--09-I07, mRNA sequence.
CF319122
Generation and analysis of 280,000 human Genome Res. 6 (9), 807-828 (1996)
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                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note="Vector: pCR4-TOPO; Site_1: ECORI; Callus was treated with ABA(20um) for 1hr. Oligo-capped mRNA was reverse transcribed and then used for PCR. mRNA was derived from rice Histone Deacetylase overexpression
                                                                                                                                                                                                                                                                                                                                                                                GI:650957
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/mol_type="mRNA"
/cultivar="Nackdong"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tissue_type="callus"
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/clone="HD--09-I07"
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Pred. No. 59;
O; Mismatches
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                         expressed sequence tags
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                                                                                                                                                                                                                                                                                                                Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                        DKFZp566B2246, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High qality sequence starts: 1
High qality sequence stops: 1
Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8889549
Other_ESTs: yb08h08.rl
Contact: Wilson RK
                                                                                                                                                  Ingolstaedter Landstr.1, D-89
Location/Qualifiers
                                                                                                                                                                                                            EST (Ottenwaelder,
Unpublished (1999)
                                                                                                                                                                                                                                                     Ottenwaelder, B., Obermaier, B.,
                                                                                                                                                                                                                                                       Mammaila; Butheria; Primates;
1 (bases 1 to 20)
Ottenwaelder
                                                                                                                                                                                                                                                                                                                                                                                                                        AL038460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                              Contact: MIPS
                                                                                                                                                                                                                                         Wiemann, S
                                                                                                                                                                                                                                                                                                                                                              AL038460.1 GI:49682131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Seq primer: -21m13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
/organism="Homo sapiens"
/mol type="mRNA"
/db xref="texon:9606"
/clone="pKrZp566B2246"
/tissue_type="kidney"
/dev stage="fetal"
/lab_host="X1-zblue"
/clone_lib="566 (synonym: hfkd2)"
/note="Vector: pAMP1; Site_1: NotI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="GDB:491520"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /db_xref="taxon:9606"
/clone="IMAGE:70623"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.1%;
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Pred. No. 69;
                                                                                                                                                                  D-85764 Neuherberg,
                                                                                                                                                                                                                                                                                                                                                                                          20 bp .....
onym: hfkd2) Homo
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                                                                                                                                                                                                                                                                                    Catarrhini;
                                                                                                                                                                                                                                                                                                  Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                       Mewes, H.W.,
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                                                                                                                                                                                                                                                                                    Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                                                      linear EST 06-JUL-2004 sapiens cDNA clone
                                                                                                                                                                                                                                                          Gassenhuber, J. and
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                                                                                                                                                                  Germany
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NotI; Site_2: SalI"

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RESULT 117
CF339443
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Best Local S
Matches 17
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JOURNAL
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17; Conserv
Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 20)
Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          _
                                                                                                                                 CF339443 20 bp mRNA linear EST 18-AUG-2003 RCL1--04-003.gl Regenerated callus lambda phage cDNA library (RCL1) Cryza sativa (japonica cultivar-group) cDNA clone RCL1--04-003,
                                                                                                                                                                                                                                                                                                                          l Similarity
17; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University Yongin, Kyeonggi, Korea Tel: 82 31 330 6193 Fax: 82 31 321 6355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7LEAF--01-D19.b1 sativa (japonica
                                                                                   CF339443.1 GI:33827271
                                                                                                                     mRNA sequence.
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                                                                                                                                                                                                                                                                                       АЛТТАЛАЛАЛАЛАЛАСТАЛ 1535
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                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /mol_type="mRNA"
/cultivar="Nackdong"
/db_xref="ttaxon:39947"
/clone="7LEAF"-01-D19"
/tissue_type="leaf"
                                                                                                                                                                                                                                                                                                                                                                                                       /dev_stage="7 days after germination"
/lab host="E.coli DHIOB"
/clone_lib="Rice leaf plasmid cDNA library II (7LEAF)"
/clone_tib="Rice leaf plasmid cDNA library II (7LEAF)"
/note="Vector: pCR4-TOPO; Site 1: EccRI; mRNA was capped
with oligoribonucleotides and then used as templates for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               organism="Oryza sativa (japonica cultivar-group)"
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Pred. No. 63;
0; Mismatches
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Pred. No. 63;
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AJ661013
                                                                                                                                                                                                                                                                        embryo development in pigs and Unpublished (2004)
                                                                                                                                                                                                                                                                                                       Anderson, S.I., Finlayson, H.A. and Archibald, A.L. Development of cDNA and EST resources for studying reproduction and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yongin, Kyeonggi, Ko
Tel: 82 31 330 6193
Fax: 82 31 321 6355
                                                                                                                                                                                                                                        Contact: Anderson SI
Genomics and Bioinformatics
                                                                                                                                                                                                                                                                                                                                                                                             Sus scrofa
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                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /mol_type="mRNA"
/cultivar="Nackdong"
/db_xref="taxon:39947"
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                                    organism="Sus scrofa"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tissue_type="callus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        clone="RCL1--04-003"
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85.0%;
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Pred. No. 63;
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1 (bases 1 to 20)
Kim, J.S., Jun, K.M., Cheong, P.J., Kim, M.J., Lee, T.H., Song, S.I., Kim, J.K., Kim, Y.-K. and Nahm, B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ehrhartoideae; Oryzeae; Oryza.
                                                                                                           /note="Vector: pBluescript SK(+); Site_1: SstI; Site_2: XhoI; CDNA was inserted into lamda Uni-ZAP XR vector at 5; end with SstI and 3; end with XhoI site. Callus was induced on 2N6 media for 30 days and cultured for 36hrs on
                                                                                                                                                                                                                                                                      /clone lib="Regenerated callus lambda phage cDNA library
(RCL1)"
                                                                                                                                                                                                                                                                                                                                              dev_stage="proliferated callus on 2N6 media for 30 days"
lab_host="E.coli SOLR"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (japonica cultivar-group)"
Length 20;
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Indels Gaps

0

AJ661013 CSEQRANO9 Sus scrofa cDNA clone C0000935_H04, mRNA EST 28-JUN-2004

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Butheria; Cetartiodactyla; Suina; Suidae; 1 (bases 1 to 21) Euteleostomi; Sus.

Roslin Institute
Roslin, Midlothian, EH25 9PS, UNITED KINGDOM
Single pass sequencing. Bases called and trimmed with phred
Single pass sequencing. Bases called and trimmed with the -minscore 20
v0.020425.c. Vector identified by cross_match with the -minscore 20
and -minmatch 12 options. Vector:pBlueScriptII(KS+) R. Site 1:
ECORI R. Site 2: NotI Description: Normalised library constructed
from pooled tissue from day 30 placentas. Clones available from UK
Centre for Functional Genomics in Farm Animals, Roslin Institite,
Roslin, Midlothian, UK, EH25 9PS, www.arkgenomics.org.

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RESULT 120
AL038627
LOCUS
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Best Local
                                AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            source
                                                                                                                                                                                                                                                                                                                         1518 ТТАЛАЛАЛАЛАЛАСТАЛАЛ 1537
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                                                                                                                                                                                       AL038627
DKFZp566H2046_r1 566 (synonym:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ALO38582 21 bp mRNA linear EST 06-JUI DKFZp566F0946 rl 566 (synonym: hfkd2) Homo sapiens cDNA clone DKFZp566F0946, mRNA sequence.
ALO38582
                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 21)
                                                                                                                                                            DKFZp566H2046, mRNA sequence.
AL038627
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Unpublished (1999)
                              Ottenwaelder, B., Obermaier, B.,
                                                                                                  Homo sapiens
                                                                                                                                              AL038627.1 GI:49682173
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Location/Qualifiers
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                                                                                                               lomo sapiens (human)
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(Ottenwaelder,
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                                                                                                                                                                                                                                                                                                                                                                                                                 /mol type="mRNA"
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/dev_stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /tissue_type="placenta"
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/clone="Yetcor: pBlueScriptII(KS+); Site_1: EcoRI; Site_2:
NotI; Single pass sequencing. Normalised library
constructed from pooled tissue from day 30 placentas."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         organism="Homo sapiens"
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et al.)
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Pred. No. 6
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nym: hfkd2) Homo
                                Mewes, H.W., Gassenhuber, J. and
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RESULT 122
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JOURNAL
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Best Local
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sequence.
CF330439
                         CF330439 21 bp mRNA linear EST 18-AUG-2003
NACL--06-C12.b1 Rice callus plasmid cDNA library (NACL) Oryza
sativa (japonica cultivar-group) cDNA clone NACL--06-C12, mRNA
                                                                                                                                       1 TCAAAAAAAAAAAAAAAAA 20
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Contact: MIPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
1 (bases 1 to 21)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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Location/Qualifiers
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Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ottenwaelder, B., Obermaier, B.,
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/mol type="mRNA"
/db_xref="taxon:9606"
/clone="bKr2p566H2046"
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/dev_stage="fetal"
/lab_host="X1-2blue"
/clone_lib="566 (synonym: hfkd2)"
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                                                                                                                                                                                                                                                         /mol_type="mrNA"
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/note="Vector: pAMP1; Site_1: NotI;
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85.0%; Pred. No. 66;
tive 0; Mismatches
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Primates;
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Pred. No. 66;
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CDNA clone
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JOURNAL COMMENT
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Oryza sativa (japonica cultivar-group)
Oryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
1 (bases 1 to 21)
Sim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Lee,T.H., Shin,Y.C.,
Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H.
Large-scale Sequencing Analysis of Rice ESTs
Unpublished (2003)
                               UMR BiO3P, BP 35327, F-35653 Le Rheu Cedex France
Tel: +33.2.23.48.51.65
Fax: +33.2.23.48.51.50
Risk of contamination by bacterial sequences from obligatory
(Buchnera) or facultative endosymbionts. These sequences were
obtained in the frame of the International Consortium of Aphic
Genomics in collaboration with Genoscope
                                                                                                                                                                                       Unpublished (2004)
Contact: D. Tagu
                                                                                                                                                                                                                                             Hunter,W., Martinez-Torres,D., Rahbe,Y., Sabater-Munoz,B., Stern,D., Tagu,D. and Wincker,P.
An expressed sequence tags database for the pea aphid Acyrthosiphon
                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Paraneoptera; Hemiptera; Sternorrhyncha; Aphidiformes;
Aphidoidea; Aphididae; Macrosiphini; Acyrthosiphon.
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CN763587.1 GI:47537510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mRNA sequence.
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ID0AAA7BH12RM1
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EST.
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Acyrthosiphon pisum
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Tel: 82 31 330 6193
Fax: 82 31 321 6355
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FORWARD: CAGGAAACAGCTATGACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /db_xref="taxon:39947"
/clone="NACL--06-C12"
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/clone lib="Rrice callus plasmid cDNA library (NACL)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was capped with oligoribonucleotides and then used as templates for RT-PCR."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Oryza sativa (japonica cultivar-group)"
/mol_type="mRNA"
/cultivar="Nackdong"
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85.0%;
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ApMS Acyrthosiphon pisum
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Pred. No. 66;
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                                                                                                                                                                                                                              Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0156 row: F column: 13
Seq primer: CGTTGTAAAACGACGGCCAGT
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                                                                                                                                                       Class: plasmid ends
High quality sequence stop:
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (2000)
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/note="Vector: pB
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/lab_host="XL1-Blue"
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/cultivar="developmentstage"
                        /db_xref="taxon:10090"
                                            /mol_type="genomic DNA"
/strain="C57BL/6J"
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clone="UUGC1M0156F13"
                                                                                                organism="Mus musculus"
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/lab_host="E. Coli strain XL10-Gold, T1-resistant,

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JOURNAL COMMENT
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AUTHORS
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ORGANISM
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AZ625662/c
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                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error:
Plate: 0465 row: C column: 23
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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Unpublished (2000)
Contact: Robert B. Weiss
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Location/Qualifiers
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GSS.
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                                                                                                                                                                                                                                                                                                                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    University of Utah Genome Center University of Utah
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(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114 gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XI10-Gold (Stratagene) cells
and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
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                                                                                            /mol_type="genomic DNA"
/strain="C57BL/6J"
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/note="Vector: PWD42nv; Purified genomic DNA |
                                                                                                                                                      organism="Mus musculus"
                             'sex="Male"
                                                  clone="UUGC1M0465C23"
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                                                                               xref="taxon:10090"
host="E. Coli strain XL10-Gold, T1-resistant,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.2;
Pred. No. 66;
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Oryza sativa (japonica cultivar-group)
Cryza sativa (japonica cultivar-group)
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CF300172 23 bp mRNA linear EST 15-AUG-2003 7LEAF--04-H15.b1 Rice leaf plasmid cDNA library II (7LEAF) Oryza sativa (japonica cultivar-group) cDNA clone 7LEAF--04-H15, mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Nahm B.H.
Genomics and Genetics Institute, GreenGene Biotech Inc.; Division of Bioscience and Bioinformatics, MyongJi University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kim,J.S., Jun,K.M., Cheong,P.J., Kim,M.J., Le Song,S.I., Kim,J.K., Kim,Y.-K. and Nahm,B.H. Large-scale Sequencing Analysis of Rice ESTs Unpublished (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                              Yongin, Kyeonggi, Korea
Tel: 82 31 330 6193
Fax: 82 31 321 6355
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                                                                                                                                                                                                                                                                                                                                                                     bhnahm@ggbio.com, bhnahm@bio.myongji.ac.kr
Location/Qualifiers
/dev_stage="7 days after germination"
/lab_host="E_coli_DH10B"
/clone_lib="Rice leaf plasmid cDNA library II (7LEAF)"
/note="Vector: pCR4-TOPO; Site_1: EcoRI; mRNA was cappe
with oligoribonucleotides and then used as templates fc
RT-PCR."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                                                                /organism="Oryza sativa (japonica cultivar-group)"
|mol_type="mRNA"
|culTivar="Nackdong"
|/db_xref="taxon:39947"
|/clone="7LEAF--04-H15"
|/tissue_type="leaf"
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Pred. No. 66;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shin, Y.C.,
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Query Match Best Local Similarity

1.1%;

Score 15.2; Pred. No. 72

DB

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RESULT 127
AW245585/c
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MI Unpublished (1999)

Other_ESTs: 2822740.5prime
Contact: Robert Strausberg, Ph.D.

Email: cgapbs r@mail.nih.gov

Email: consortium (LIMI) DNA Sequencing by: Berkeley MGC sequencing by

Berkeley MGC sequence Can be found through the I.M.A.G.E. Consortium/LIMI at:

www-bio.llnl.gov/bbrp/image/hmage.html Base Calling / Quality

Scores: PHRED from University of Washingtion Genome Center:

Frimming: cross match from University of Washingtion Genome Center:

FRAPP suite. Poly-T Identification: patMatch.pl from Berkeley

Drosophila Genome.washington.edu Low Quality Sequence: 6 contiguous

PHRED high quality bases following vector sequence. Very Low

Quality Sequence: Trace file contained 15 contiguous distinct peaks

following vector sequence. Polyadenylation: Based upon the presence

of a Xhol site followed by a run of 14 or more T residues at the

beginning of the sequence, this cDNA insert was polyadenylated.

Flocation/Onalificats

Location/Onalificats

Location/Onalificats

Location/Onalificats
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National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1. (bases 1 to 15)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AW245585.1 GI:6588578
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mRNA sequence.
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                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                     /clone lib="NIH_MGC_7"
/note="Organ: lung; Vector: pOTB7; Site 1: XhoI; Site 2: EcoRI; cDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit
                                                                                                                                                                                                                                                                           California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone="IMAGE:2822740"
/tissue_type="small cell carcinoma"
/cell_line="MGC3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'lab_host="DH10B (phage-resistant)"
                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                               1.1%; Score 15;
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KEYWORDS
Search completed: November Job time : 3 secs
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ORGANISM
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LOCUS
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Best Local :
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RESULT 128

1518 TTAAAAAAAAAAAA 1532

4 TTAAAAAAAAAAAAG

18

2, 2004, 12:50:35

15;

Conservative

Mismatches

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Gaps

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AZ774536 19 bp DNA linear GSS 10 2M0004P01F Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC2M0004P01 F, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Plate: 0004 row: P column: 01
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            High quality sequence stop: 19.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert B. Weiss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus
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Fax: 801 585 7177
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                                                                               (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sex="Male"
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100.0%;
1.1%; Score 15;
100.0%; Pred. No.
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                          Length 19;
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                                                                                                                                                                                     Shannon, M., Gu, Y. and Nguyen, C.T. Four human zinc-finger-containing proteins: mdz3, mdz4, mdz7
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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RESULT 7
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Sequence 4 from Patent WO03072818.
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AX825106.1 GI:39750835
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Similarity 87.0%;
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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synthetic construct
artificial sequences.
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Vaisberg, E.A., Adams, C.L., Sabry, J.H. and Crompton, A.M.
Database system including computer code for predictive cellular
                                                                                                                                                                                                                                                                                                           Sequence 1 from patent US AR431307
                                                                                                                                                                     Patent: US 6651008-A 1 18-NOV-2003;
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                                                                                                         /organism="unknown"
/mol_type="genomic DNA"
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/mod_base=OTHER
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/mod_base=OTHER
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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AX825103
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Penny,L. and Galvin,M.
Genotyping the human UDP-glucuronosyltransferase 1 (UGT1) gene
Patent: US 6479236-A 100 12-NOV-2002;
Location/Qualifiers
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mol_type="unassigned DNA"

db_xref="taxon:32630"
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90.0%; Pred. No. 53;
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Sequence 2 from Patent WO03072818.
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Sequence 3 from Patent WO03072818.
AX825105
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Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucleic acids
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Degussa Bioactives GmbH (DE)
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                                                      /note="LNA-T (Locked
/mod_base=OTHER
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/mal_type="unassigned DNA"
/mal_type="taxon:32630"
/db_xref="taxon:32630"
/noTe="Beschreibung der_kuenstl
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Chetverin,A.B. and Kramer,F.R.
Oligonucleotide arrays and their use for sorting,
sequencing, and manipulating nucleic acids
patent: US 6322971-A 6 27-NOV-2001;
Location/Qualifiers
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                                                                                                                                     Patent: WO 8404538-A 24 22-NOV-1984;
Location/Qualifiers
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Sequence 6 from patent US
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AR294636.1
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Sequence
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Cohen,D., Chumakov,I. and Blumenfeld,M.
Biallelic markers for use in constructing
disequilibrium map of the human genome
Patent: US 6537751-A 6371 25-MAR-2003;
Location/Qualifiers
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1 (bases 1 to 23)

Edens,L., Russell,S.W., Visser,C.

Improvements in the expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Reverse complement, A06442
Unclassified.
1 (bases 1 to 21)
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UNILEVER NV; UNILEVER PLC
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6371 from patent US
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
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Oligonucleotide repeat arrays
Patent: US 5981195-A 55 09-NOV-1999;
Location/Qualifiers
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                                                                                                                     Matson, R.S., Coassin, P.J., Rampal, J.B. Oligonucleotide repeat arrays Patent: US 5981185-A 68 09-NOV-1999; Location/Qualifiers
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Sequence 68 from patent US
AR084579
AR084579.1 GI:10011350
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Matson,R.S., Coassin,P.J., Rampal,J.B.
Oligonucleocide repeat arrays
Patent: US 5981185-A 67 09-NOV-1999;
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Sequence 3 from Patent WO2004055153.
CQ830491
CQ830491.1 GI:50250811
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Matson,R.S., Coassin,P.J., Rampal,J.B.
Oligonucleotide repeat arrays
Patent: US 5981185-A 71 09-NOV-1999,
Location/Qualifiers
Schluesener,H. and Wendel,H.P. Devices coated with substances that mediate biological material Patent: WO 2004055153-A 3 01-JUL-2004; Eberhard-Karls-Universitaet Tuebingen (DE)
                                                                                                                                                                                                                                                   18;
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Bergan,R. and Neckers,L.
Inhibition of protein kinase activity by aptameric action of
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AR093142
                                                                              synthetic construct synthetic construct
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                                                                   artificial sequences.
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Similarity 85.7%;
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/mol_type="unassigned
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/mol_type="unassigned DNA"
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Patent: WO 0214546-A 34 21-FEB-2002;
                                                                                                                                                                                                               Use of microbial dna sequences for the
                                                                                                                                                                                                                                                  Plasmodium falciparum (malaria parasite P. falciparum)
Plasmodium falciparum
Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium.
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1 (bases 1 to 21)
Milhausen, M.J.
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larity 85.7%; Pred. No. 7
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                                                                                                      /organism="Plasmodium
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db_xref="taxon:32630"
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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/db_xref="taxon:32630"
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/mod_base=OTHER
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/mol type="unassigned DNA"
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Sequence 20 from Patent WO03072818
AX825122
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Location/Qualifiers
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/mol type="unassigned DNA"

/mol type="unassigned DNA"

/db_xref="taxon:32630"

/note="Beschreibung der kuenstlichen

Sequenz:Capture-Oligonukleotid"
   /note="LNA-T (Locked Nucleic Acid)"
                                                                                                /note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
                                                                                                                             organism="synthetic construct"

/mol_type="unassigned DNA"

/db_xref="taxon:32630"
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/mod_base=OTHER
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/mod_base=OTHER
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mod_base=OTHER
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mod_base=OTHER
                         mod_base=OTHER
                                             note="LNA-T (Locked Nucleic Acid)"
                                                                     bound_moiety="Biotin"
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AX825138/c
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Sequence 36 from Patent W003072818.
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
                                                                                                                                               /note="LNA-T {Locked
/mod_base=OTHER
                                                                                                        /note="LNA-T (Locked Nucleic Acid)"
/mod_base=OTHER
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85.7%;
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Pred. No. 76;
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Pred. No. 76;
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BD085544
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AX825154/c
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                                        Method of comparison and detection of RNA Patent: JP 2001333800-A 1 04-DEC-2001; UNITECH CO LTD
                                                                                            Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
L. (bases 1 to 22)
                                                                                                                                                                                             BD085544
Method of comparison and
BD085544
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                                                                                                                                                  Homo sapiens (human)
                                                                                                                                                                   BD085544.1 GI:22631154
JP 2001333800-A/1.
                                                                                  Shimada, K.
                                                                                                                                         Homo sapiens
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Homo sapiens (human)
JP 2001333800-A/1
04-DEC-2001
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/mod_base=OTHER
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/mal_type="unassigned DNA"
/mb xref="taxon:32630"
/note="Beschreibung der kuenstlichen
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AR315696/c
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Sequence
AR315696
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                                                         1 (bases 1 to 20)
Vasseur,M., Blumenfeld,M., Meguenni,S. and Poddevin,B.
STAPLE AND SEMI-STAPLE OLIGONUCLEOTIDES, METHOD OF PRE
APPLICATIONS
Patent: WO 9423026-A 2 13-OCT-1994;
GENSET (FR)
                                                                                                                                                                                           20 bp
Sequence 2 from Patent WO9423026.
A40126
                                                                                                                                                                                                                                                                                                                                  16;
                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 20)
Griffals,R., Hoiseth,S.K., Zagursky,R.J., Metcalf,B.J.,
Sankaran,B. and Fletcher,L.D.
Chlamydia pneumoniae polynucleotides and uses thereof
Patent: US 6559294-A 6233 06-MAY-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N
                                                                                                                                  unidentified unclassified.
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C12Q1/68,C12N15/09,G01N33/50,C12N15/00
Method of comparison and detection of RNA amount and
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                     publication AU 6432094 941024
publication FR 2703053 940930
Location/Qualifiers
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    .20
    /organism="unidentified"

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/mol_type="genomic RNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                /mol_type="genomic DNA'
                                                                                                                                                                                   GI:2296284
                                                                                                                                                                                                                                                                                                                                                                                              organism="unknown"
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100.0%; Pred. No.
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85.7%;
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1. .22
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Pred. No. 73;
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6559294.
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                     RESULT 38
AR140558/c
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AR140280/c
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AR139961/c
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al Similarity 89.5%;
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AR139961
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Spebo, K.M., Bosselman, R.A., Suggs, S.V. and Martin, F.H.

Method for enhancing the effciency of gene transfer with stem cell
factor (SCF) polypeptide
Patent: US 6207454-A 33 27-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unknown
                                                                                                                                                                                                                                                                 Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA encoding stem cell factor Patent: US 6207417-A 33 27-MAR-2001;
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                                                                                                                                                                                                                                                     Unclassified.
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Zsebo,K.M., Bosselman,R.A., Suggs,S.V.
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                                                                                                                                                   /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned
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/db_xref="taxon:32644"
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                                                                                                                     1.1%; Score 15.8; DB 1; 89.5%; Pred. No. 1e+02;
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AX104051
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ORGANISM
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AR182885
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AR182885
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1 (bases 1) to 20)
Krieg,A.M., Davis,H.L., Wu,T. and Schorr,J.
Vectors and methods for immunization or therapeutic protocols
Patent: US 6339068-A 57 15-JAN-2002;
Location/Qualifiers
                                                                                                                                                                                                  Sequence 243 from Patent AX104051
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Stem cell factor and compositions
Patent: US 6207802-A 33 27-MAR-2001;
                                                                       Krieg, A.M., Schetter.C. and Vollmer,J.C. Immunostimulatory netleic acids Patent: WO 0122972 nc.41 05-APR-2001; UNIVERSITY OF IOWA RESEARCH FOUNDATION (US); Coley Pharmaceutical
                                                                                                                                                  synthetic construct synthetic construct
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AR140558.1
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                                                                                                                                     artificial sequences.
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ilarity 89.5%;
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organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                     /mol_type="unassigned DNA"
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AX281587
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DEFINITION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              synthetic construct synthetic construct
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                                                                                                                                                       Patent: WO 0197843-A 410 27-DEC-2001;
UNIVERSITY OF IOWA RESEARCH FOUNDATION
Location/Qualifiers
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                                                                                           /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="raxon:32630"
/noTe="Synthetic oligonucleotide-phosphodiester
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/noTe="Synthetically generated primer"
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Pred. No. 1e+02;
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Use of nucleic acids containing unmethylated CPG dinucleotide in the treatment of LPS-associated disorders

LE Patent: JP 2001513776-A 65 04-SEP-2001;

UNIVERSITY OF IOWA RESEARCH FOUNDATION

OS Artificial Sequence
PD 04-SEP-2001
PF 25-PEB-1991 JP 1998537810
PF 25-PEB-1998 JP 1998537810
PF 25-PEB-1999 UP 1998537810
PF 26-PEB-1991 UP 1998537810
PF 26-PEB-1991 UP 1998537810
PF 26-PEB-1991 UP 1998537810
PR 28-PEB-1991 UP 1998537810
PF 28-PEB-1998 UP 1998537810
PF 38-PEB-1998 UP 1998537810
PF 28-PEB-1998 
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Bratzler,R.L.

Inhibition of angiogenesis by nucleic acids Patent: WO 02053141-A 243 11-UUL-2002; Coley Pharmaceutical Group, Inc. (US)

Location/Qualifiers
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JP 2001513776-A/65.
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Use of nucleic acids containing the treatment of LPS-associated
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ilarity 89.5%;
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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l Similarity 89.5%;
17; Conservative
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Matson,R.S., Coassin,P.J., Rampal,J.B. and Caskey,C.Thomas.
Oligonucleotide repeat arrays
Patent: US 5981185-A 56 09.NOV-1999;
Location/Qualifiers
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Matson, R.S., Coassin, P.J., Rampal, J.B. and Caskey, C.Thomas.
Oligonucleotide repeat arrays
Patent: US 5981185-A 52 09-NOV-1999;
Location/Qualifiers
Schluesener,H. and Wendel,H.P. Devices coated with substances that mediate biological material Patent: WO 2004055153-A 2 01-JUL-2004; Eberhard-Karls-Universitaet Tuebingen (DE)
                                                                           synthetic construct
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Sequence 56 from patent US
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/mol_type="unassigned DNA"
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Sequence 4 from Patent WO2004055153.
CQ830492
CQ830492.1 GI:50250832
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Devices coated with substances that mediate the adhesion of biological material
Patent: WO 2004055153-A 4 01-JUL-2004;
Eberhard-Karls-Universitaet Tuebingen (DE)
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Ford, J. and Yeung, G.
Chemokine receptor obtained from a cDNA library of
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Patent: US 6472173-A 20 29-OCT-2002;
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/mol_type="unassigned DNA"
/mol type="unassigned DNA"
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Location/Qualifiers
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Ford, J. and Yeung, G.
Chemokine receptor obtained from a cDNA library of fetal
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/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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                                                  /note="LNA-T (Locked Nucleic Acid)"
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mod_base=OTHER
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Sequenz:Capture-Oligonukleotid"
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Sequenz:Capture-Oligonukleotid"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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   mod_base=OTHER
                                                               note="LNA-T (Locked Nucleic Acid)"
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Location/Qualifiers
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Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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SOURCE
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AX825115/c
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AX825113/c
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AX825113
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Location/Qualifiers
                                                                                                                                        synthetic construct
synthetic construct
artificial sequences.
                                                                                                                                                                                                            Sequence 13 from Patent AX825115
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Sequenz:Capture-Oligonukleotid"
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/db_xref="taxon:32630"
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Sequence 14 from Patent WO03072818.
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Sequenz:Capture-Oligonukleotid"
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/noTe="Beschreibung der kuenstlichen
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AX825152/c
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Query Match 1.1%;
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Sequence 15 from Patent W003072818.
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AX825153.1 GI:39750882
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Seguenz:Capture-Oligonukleotid"
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/mol type="unassigned DNA"
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/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
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AX825163/c
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/mol_type="unassigned DNA"
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Sequence 19
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131828
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Methods for identification of cancer cell surface molecules cancer specific promoters, and therapeutic uses thereof Patent: WO 03000928-A 415 03-UAN-2003;

Odin Medical A/S (DK)
Unclassified.
1 (bases 1 to 22)
Torrence,P., Silverman,R., Maitra,R. and Lesiak,K.
                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 22)
Torrence, P.F., Silverman, R.H., Maitra, R.K. and Lesiak, K. Chimeric molecules targeted to viral RNAs
Patent: US 6271369-A 19 07-AUG-2001;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                organism="unknown"
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Sequence 19 from patent US 5677289.
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169425.1 GI:2831547
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                         C12P21/02,C12P21/02//A61K31/711,(C12N5/10,C12R1:91),(C12P21/02, PC C12R1:91),
                                                                                              OS EUKATYOTE
PN JP 2002541795-A/2217
PD 10-DEC-2002
PF 11-APR-2000 JP 2000611654
PF 11-APR-1999 US 60/129390
PR 12-APR-1999 US 60/129390
PR 12-APR-1999 US 60/129390
PR 12-APR-1999 US 60/129390
PI LAWRENCE BLATT, MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN PC C12N15/09, A61K38/00, A61K48/00, A61P43/00, A61P43/00, C12N5/10, PC
                                                                                                                                                                                                                                1 (bases 1 to 17)

Blatt, L., Zwick, M., Pavco, P. and Mcswiggen, J.

Regulation of repressor genes using nucleic a

Patent: JP 2002541195-A 2217 10-DEC-2002;

RIBOZYME PHARMACEUTICALS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unclassified.

1 (bases 1 to 22)

Torrence,P., Silverman,R., Maitra,R. and Lesiak,K.

Method of cleaving specific strands of RNA and medical treatments
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Beigelman,L., Burgin,A., Beaudry,A., Karpeisky,A.,
Matulic-Adamic,J., Sweedler,D. and Zinnen,S.
Synthetic ribonucleic acids with RNAse activity
Patent: US 6528640-A 559 04-MAR-2003;
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Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6346398-A 2546 12-FBB-2002;
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559 from patent US 6528640.
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/db_xref="taxon:32644"
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                                            /organism="unknown"
/mol_type="unassigned RNA"
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Pavco, P., McSwiggen, J.A., Stinchcomb, D.T. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6566127-A 1070 20-MAY-2003;
Location/Qualifiers
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Oreochromis niloticus (Nile tilapia)
Oreochromis niloticus
Eukaryota; Metazoa; Chordata; Crania
                                                                                                                                                                                             16;
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Beigelman, L., Burgin, A.B., Beaudry, A., Karpeisky, A.,
Matulic-Adamic, J., Sweedler, D. and Zinnen, S.
Oligoribonucleotides with enzymatic activity
Patent: US 6617438-A 558 09-SEP-2003;
                                                                                                                                                                                                                                                                                                                                                               Unknown
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Similarity 94.1%; Pred. No. 1.
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/mol_type="unassigne
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/mol_type="unassigned RNA"
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  Chordata; Craniata; Vertebrata;
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A88305.1
                                                   unidentified unclassified.

1 (bases 1 to 20)
Brysch, W.D. and Schlingensiepen, K.D. An antisense oligonucleotide preparation Patent: EP 0856579-A 453 05-AUG-1998; BIOGNOSTIK GES (DE)
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Brysch, W. and Schlingensiepen, K.

Brysch, W. and Schlingensiepen, K.

ANTISENSE OLIGONUCLEOTIDE PREPARATION METHOD Patent: WO 9933904-A 453 06-AUG-1998;

BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              unidentified unidentified
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Sequence 453 from Patent
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                                                                                                                                               unidentified
                                                                                                                                                                       A90272.1 GI:6738786
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  unclassified.
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ilarity 94.1%;
Conservative
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/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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/mol_type="unassigned DNA"
/db_xref="taxon:8128"
                                        Location/Qualifiers
/organism="unidentified"
/mol_type="unassigned DNA"
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Pred. No. 1.4e+02;
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Pred. No. 1.3e+02;
0; Mismatches 1;
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E 1 (bases 1 to .20)

S Schlingensiepen, K.H. and Brysch, W.

An antisense oligonuclectide preparation method

Patent: JP 2001511000-A 453 07-AUG-2001;

BICCROSTIK GESELLSCHAFT FUR BICMOLEKULARE DIAGNOSTIK MBH

OS UNKNOWN

PN JP 2001511000-A/453

PD 07-AUG-2001

PF 30-JAN-1998 JP 1998532533

PF 30-JAN-1997 EP 97101531.8

PR 31-JAN-1997 EP 97101531.8

PR 31-JAN-1997 EP 97101531.7

CC An antisense oligonuclectide preparation method FH Ke
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AX356277
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An antisense oligonucleotide
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Plant regulatory sequences for selective control of gene expression
Patent: WO 0200905-A 71 03-VAN-2002;
Monsanto Technology LLC (US)
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Pred. No. 1.3e+02;
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AR112167/c
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Best Local Similarity
Matches 16; Conserv
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AR112167 20 bp
Sequence 56 from patent US 6130041.
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Acton,S.Laurene.
Intronic and polymorphic SR-BI nucleic acids
Patent: US 5998141-A 56 07-DEC-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                       Sequence 56 from patent US
AR092032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genotyping reagents, kits and methods of use thereof Patent: WO 0129262-A 3390 26-APR-2001; Orchid BioSciences, Inc. (US)
                                                                                                                                                                                                                                                                                                                 Unknown
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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                                                                                                                                                                                       /mol_type="unassigned DNA"
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Pred. No. 1.2e+02;
0; Mismatches 1;
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Pred. No. 1.3e+02;
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Sequence
AR149209
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 20)

Galvin,M., Miller,A., Penny,L. and Riedy,M.
Genotype determination of human UDP-glucuronosyl transferase 2B4 (UGT2B4), 2B7 (UGT2B7) and 2B15 (UGT2B15) genes
Patent: JP 2002521067-A 90 16-JUL-2002;

AXYS PHARMACEUTICALS INC
OS Homo sapiens (human)
                                                                                                                                                                              20 bp DNA linear PAT 17-JU Genotype determination of human UDP-glucuronosyl transferase (UGT2B4), 2B7 (UGT2B7) and 2B15 (UGT2B15) genes.
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Acton, S.L. and Ordo
Human intronic and
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Acton,S.Laurene.
Human_intronic and polymorphic SR-BI nucleic acids and uses
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                                                                                                                                                       BD229218.1 GI:33038988 JP 2002521067-A/90.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent: US 6228581-A 56 08-MAY-2001;
Location/Qualifiers
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AR112167.1
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                                                                                                                            Homo sapiens
                                                                                                                                       Homo sapiens (human)
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/mol_type="unassigned DNA"
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85.0%;
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AR349470
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Matches 17; Conserv
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Best Local Similarity 85.0%;
Matches 17; Conservative
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PD 16-UUL-1909 JP 2000562558

PF 22-UUL-1999 JP 2000562558

PR 28-UUL-1998 US 60/094391

PI MARGARET GALVIN, ANDREW MILLER, LAURA PENNY, MICHAEL RIEDY F

C12N15/09, C12N15/09, C12M1/00, C12Q1/68, C12N15/00, C12N15/00 CC

Genotype determination of human UDP-glucuronosyl transferase

284 (UCT2B4),

284 (UCT2B4) genes
                                     Gal. (bases 1 to 20)
Gal. (h.m., miller, A., Penny, L. and Riedy, M.
Genotyping the human UDP-glucuronosyltransferase
Patent: US 6586175-A 92 01-UUL-2003;
Location/Qualifiers
                                                                                                                    Unknown
                                                                                                                                                                        AR349470
Sequence 92 from patent US
AR349470
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Aizawa,A., Kawakami,A. and Kondo,T.
Testis-specific gene
Patent: US 6555670-A 4 29-APR-2003;
Location/Qualifiers
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Sequence 4 from patent US
AR309844
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
             organism="unknown"
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_type="genomic DNA"
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Pred. No. 1.5e+02;
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Pred. No. 1.5e+02;
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AX404077
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Patent: WO 0183749-A 175 08-NOV-2001;
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                                                                                                                                                            Livestock Improvement Association of Gunma University (JP)
                                                                                                                                                                                   Testis-specific gene
Patent: EP 1195382-A 4 10-APR-2002;
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AX298541
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                                                                                                                                                                                                                                   artificial sequences.
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                                                                                          /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned |
/db_xref="taxon:10095"
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                                                                                                                                             Location/Qualifiers
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Pred. No. 1.5e+02;
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                                              Score 15.2; DB 1
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BD143136.1 GI:27848894
JP 2002112777-A/3
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Sequence 9 from Patent WO03016562.
AX770111
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Sequence 5708 from Patent
AX488408
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/db_xref="taxon:32630"
/note="Primer"
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/db_xref="taxon:5476"
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Aizawa,A., Kawakami,A. and Kondo,T.
Novel testis-specific gene
Patent: JP 2002112777-A 3 16-APR-2002;
KACHIKU KAIRYO JIGYODAN, PRESIDENT OF GUNMA UNIVERSITY
                                                                                                                                                                                                                                                                                                                                                                          Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucle Patent: WO 03072818-A 4 04-SEP-2003; Degussa Bioactives GmbH_(DE)
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Sequence 4 from Patent WO03072818.
AX825106
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Similarity 85.0%; Pred. No. 1.5e+02
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AKIRA AIZAWA, AKIKO KAWAKAMI, TOSHIHIKO
C12N15/09, C07K14/47, C12N15/00
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JP 2002112777-A/3
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/mol type="unassigned DNA"
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/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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                                                                                                               /note="LNA-T (Locked
/mod_base=OTHER
                                                                                                                                                             /mod
    /note="LNA-T (Locked Nucleic Acid)'
                          mod_base=OTHER
                                              /note="LNA-T (Locked
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A20531 21 bp DNA lin oligonucleotide for the mutagenesis of SAT216.
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synthetic construct
artificial sequences.
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1 (bases 1 to 21)
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Location/Qualifiers
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Location/Qualifiers
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ilarity 85.0%;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon;32630"
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Pred. No. 1.4e+02;
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Pred. No. 1.4e+02;
0; Mismatches 3;
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Pred. No. 1.4e+02;
0; Mismatches 3;
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Best Local S
Matches 17
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AR143626
AR143626.1 GI:15104912
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Location/Qualifiers
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A20531.1 GI:583363
              Patent: US 6204435-A 11 20-MAR-2001;
Location/Qualifiers
                                                               Stamp, L.M.
                                                                     1 (bases 1 to 21)
Feitelson,J.S., Schnepf,H.Ernest., Narva,K.E., Stockhoff,B.A., Schmeits,J., Loewer,D., Dullum,C.Joseph., Muller-Cohn,J. and Stamp 1. M
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                                                  Pesticidal toxins and nucleotide sequences which encode these
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Matches 17; Conserv
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I (bases) to 21)

Babba, T. and Shimada, H.

NEW RICE PLANT STARCH-BRANCHED ENZYMIC GENE
PAtent: JP 1994261767-A $ 20-SEP-1994;

MITSUI GIYOUSAI SHOKUBUTSU BIO KENKYUSHO:KK

OS None

OC Artificial sequences.

PN JP 1994261767-A/5

PD 20-SEP-1994

PF 22-OCT-1993 JP 1993265171

PR 29-OCT-1993 JP 1993265171

PR 29-OCT-1992 JP 92P 291719

PI BABA TADASHI, SHIMADA HIROAKI

PC C12N15/54,A01H5/00,C12N5/10,C12P19/16//A2

BITANDAGHI, SHIMADA HIROAKI

PC topology: Linear;
FH Key Location/Qualifiers

FH Key / Organism='Artificial seq
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E08187.1 GI:2176308
JP 1994261767-A/5.
unidentified
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AR157200
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17; Conservative
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Feitelson, J. Schnepf, H. Ernest, Narva, K.E., Stockhoff, B.A., Schmeits, J., Loewer, D., Dullum, C. Joseph., Muller-Cohn, J., Stan Morrill, G. and Finstad-Lee, S.
Pesticidal toxins and nucleotide sequences which encode these
                                                                                                                                                                                                                                                                                                           unidentified
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Location/Qualifiers
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pri994261767-A/5
20-SEP-1994
22-OCT-1993 JP 1993265171
29-OCT-1992 JP 92P 291719
BABA TADASHI, SHIMADA HIROAKI
ELIN15/54,AO1H5/00,Cl2N5/10,Cl2P19/16//A23L1/10,Cl2N9/10;
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/mol_type="unassigned DNA"
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   organism='Artificial sequences'.
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ACCESSION
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AR437180
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Location/Qualifiers
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synthetic construct
artificial sequences.
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Feitelson, J.S., Schnepf, H.E., Narva, K.E., Stockhoff, B.A., Schmeits, J., Loewer, D., Dullum, C.J., Muller-Cohn, J., Stamp, L., Morrill, G. and Finstad-Lee, S.
Pesticidal toxins and nucleotide sequences which encode these
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Similarity 85.0%; Pred. No. 1.4e+02;
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/mol_type="genomic DNA"
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                                                                        Sequenz: Capture-Oligonukleotid"
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/mb = "Beschreibung der kuenstlichen
/note="LNA-T (Locked Nucleic Acid)"
/mod_base=OTHER
                                            /bound_moiety="Biotin"
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85.0%; Pred. No. 1.4e+02;
tive 0; Mismatches 3
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AX825120/c
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AX825120
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ilarity 85.0%;
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/mod_base=OTHER
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/db_xref="taxon:32630"
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/mod_base=OTHER
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/mod_base=OTHER
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                       Score 15.2; DB 1;
Pred. No. 1.4e+02;
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          Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucleic acids Patent: WO 03072818-A 33 04-SEP-2003; Degussa Bioactives GmbH (DE)
Location/Qualifiers
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Sequence 33 from Patent WO03072818.
AX825135
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/mod_base=OTHER
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/mod_base=OTHER
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Sequenz:Capture-Oligonukleotid"
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/mol_type="unassigned DNA"
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Pred. No. 1.4e+02;
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Sequence 34 from Patent WO03072818.
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17; Conserv
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/ppe="taxon:32630"
/note="Beschreibung der kuenstlichen
seguenz:Capture-Oligonukleotid"
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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                                           note="LNA-T (Locked Nucleic Acid)"
                                                                      mod_base=OTHER
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AX825136/c
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AX825137/c
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Sequence 34 from Patent W003072818.
AX825136
             AX825137 21 bp
Sequence 35 from Patent W003072818.
AX825137 GI:39750866
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Location/Qualifiers
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Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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/mod_base=OTHER
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/db_xref="taxon:32630"
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mod_base=OTHER
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Length Indels

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RESULT 105
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Whiches 17; Conservative
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                                                                                                                                                                                                                                                                                synthetic construct synthetic construct
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Sequence 53 from Patent WO03072818.
AX825155
AX825155.1 GI:39750884
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Sequenz:Capture-Oligonukleotid"
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/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
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                             note="LNA-T (Locked Nucleic Acid)"
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21 bp
Sequence 57 from Patent WO03072818.
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Location/Qualifiers
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/mod_base=OTHER
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/mod_base=OTHER
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/mol_type="unassigned DNA"
/mol_type="unassigned DNA"
/db xref="taxon:32630"
/noFe="Beschreibung der kuenstlichen
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/mod_base=OTHER
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B5.0%; Pred. No. 1.4e+02;
vative 0; Mismatches 3;
             1.1%; Score 15.2; DB 1;
85.0%; Pred. No. 1.4e+02;
Mismatches
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                                DB 1; Length 21;
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BD023735
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AUTHORS
T PD 2001506136-A/1
PD 15-MAY-2001
PF 29-DEC-1997 .To ...
PASCOMYCOTA; Saccharomycotina; Saccharomycetes;
Ascomycota, Saccharomycotina; Saccharomycetes;
Le Scanciales, Saccharomycotina; Saccharomycetes;
Le Compared to the second terms of the second terms.
Le Compared to the second terms of the second term
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         source
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PI JAMES L SCHWEITS, DAVID LOEWER, GEORGE SCHWAB,

PI CHARLES JOSEPH DULLUM,

PI JUDY MULLER COHN, LISA STAMP

PC C12N15/32,C07K.14/325,C12Q1/68,A01N63/00,C12N15/82

Strandedness: Single;

CC Topology: Linear;

CC Topology: Linear;

FH Key

FH Key

1. 21

FT source /organism='Unidentified'.
                                                                                                                                                                                                        BD023735.1 GI:22564958
JP 2001506136-A/1.

Eremothecium gossypii (Ashbya gossypii)

Eremothecium gossypii

Eremothecium gossypii

Eukaryota; Fung1; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Eremothecium.
                                                                                                                                                                                                                                                                                                                                               BD023735
Beta-galactosidase having
BD023735
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Peitelson, J.S., Schnepf, E.H.,
Schmeits, J.L., Loewer, D., Schw
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BD008680.1 GI:18637053
JP 2001502919-A/8.
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JP 2001502919-A/8
06-MAR-2001
30-OCT-1997 JP 1998520788
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pesticidal toxins and nucleotide sequences which encode these
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Pred. No. 1.4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
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AR187059/c
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                              /mol_type="unassigned
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1511 CTGTTAATTAAAAAAAAAA 1530
                                           1 (bases 1 to 17)
Pavco,P., McSwiggen,J., Stinchcomb,D. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 2547 12-FEB-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genotyping reagents, kits and methods
Patent: WO 0129262-A 2755 26-APR-2001,
Orchid BioSciences, Inc. (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             artificial sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                1.1%; Score 15.2; DB 1; Similarity 85.0%; Pred. No. 1.2e+02
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Strandedness: Single;
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llarity 85.0%;
Conservative
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/mol_type="genomic DNA"
/db_xref="taxon:33169"
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            organism="unknown"
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/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
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Pred. No. 1.4e+02;
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AR323670/c
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AR323669/c
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Best Local Similarity 100.0%; Pred. No. 1.9e+02;
Matches 15; Conservative 0; Mismatches 0;
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Sequence 2548 from patent
AR187060
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Sequence 1071 from patent
AR323669
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1 (bases 1 to 17)

Paver (bases 1 to 17)

Paver (bases 1 to 17)

Paver (bases 1 to 17)

Rethod and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6346398 A 2548 12-FEB-2002;
          AR323670
Sequence 1072 from patent
AR323670
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Pavco, P., McSwiggen, J.A., Stinchcomb, D.T. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor recepts
Patent: US 6566127-A 1071 20-MAY-2003;
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AR323670.1
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                                                                                                                                                                                         /organism="unknown"
/mol_type="unassigned RNA"
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Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Pavco, P., McSwiggen, J.A., Stinchcomb, D.T. and Escobedo, J
Method and reagent for the treatment of diseases or cond
related to levels of vascular endothelial growth factor
Patent: US 6566127-A 1072 20-MAY-2003;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 169 from Patent
AX753822
AX753822.1 GI:32166519
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Human angiomotin-like protein 1
Patent: WO 03037931-A 168 08-MAY-2003;
Amersham Biosciences SV Corp. (US)
Location/Qualifiers
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                                Human angiomotin-like protein 1
Patent: WO 03037931-A 169 08-MAY-2003;
Amersham Biosciences SV Corp. (US)
Location/Qualifiers
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned RNA"
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organism="Homo sapiens"/
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AR102020/c
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Sequence 170 from Patent WO03037931.
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Sequence 18 from patent US 6083731.
AR102020
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Croteau, R. Bruce., Lupien, S. Lee. and Karp, F.
Recombinant materials and methods for the phydroxylases
                                                                                                                                                                                                                                             Unknown
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Patent: WO 03037931-A 170 08-MAY-2003;
Amersham Biosciences SV Corp. (US)
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae, Homo.
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                                                                                                                                                                                                                                  Unclassified.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Pred. No. 1.7e+02;
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AR352460/c
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Sequence 6
AR352460
AR352460.1
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1241 CCTCATCTTTGTTTT 1255
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                                            1 (bases 1 to 20)
Hideki,K. and Senshu,U.
Method for analyzing DNA fragment
Patent: JP 1999196874-A 9 27-JUL-1999;
                                                                                                                                                      E28098
E28098.1 GI:13018323
JP 1999196874-A/9,
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Kong, Y., Chung, J.-Y., Bahk, Y.Y., Kang, S.-Y. and Cho, S.-Y.
Recombinant antigen of Taenia solium metacestodes
Patent: US 6589752-A 6 08-JUL-2003;
Location/Qualifiers
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Croteau, R. Bruce., Lupien, S. Lee. and Karp, F.
Recombinant materials and methods for produced
                                                                                                                                                                                                       Method for analyzing DNA fragment.
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Unidentified 
JP 1999196874-A/9
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PP 14-JAN-1998 JP 1998005399
PR HIDEKI KAMIBARA, SENSHU UEMATSU
PI C12N15/09, C12Q1/68, G01N27/447, C12N15/00, G01N27/26
Strandedness: Single;
CC Topology: Linear;
FH Key Location/Qualifiers
FH Key 1.20
FT Bource 1.20
/organism='Unidentified'.
                                                                                                                                                                                                                                                                                                                                                                                                                        Unclassified.

1 (bases 1 to 20)

Cohen, D., Chumakov, I. and Blumenfeld, M.
Biallelic markers for use in constructing
disequilibrium map of the human genome
Patent: US 6537751-A 6435 25-MAR-2003;
Location/Qualifiers
                                                                                                                                                                AR307942
Sequence 153 from patent
AR307942
AR307942.1 GI:31698698
                                      Antisense modulation of raidd express
Patent: US 6551826-A 153 22-APR-2003;
Location/Qualifiers
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Sequence 6435 from patent
AR294700
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/mol_type="genomic_DNA"
/db_xref="taxon:32644"
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Location/Qualifiers
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              organism="unknown"
mol_type="genomic DNA"
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tive 1; Mismatches 2;
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100.0%; Pred. No. 1.6e+02;
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AX048446/c
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AX048446
AX048446.1
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Sequence 1012 from Patent
AX804844
                                                                                                                                                                                                     Lie,Y., Slettan,A., Hoeyum,M. and Lingaas,F. Verification of food origin based on nucleic
                                                                                                                                                                                                                                              Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Buteleostei; Neoteleostei; Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Labroidei; Cichlidae; Oreochromis.
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                                                                                                                                                                              Patent: WO 03060160-A 1012 24-JUL-2003;
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                                                                                          /organism="Oreochromis niloticus"
/mol_type="unassigned DNA"
/db_xref="taxon:8128"
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/db_xref="taxon:32630"
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                                                                                                                                    AR058305
Sequence 3
AR058305
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1 (bases 1 t
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AR034896
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Chatelain, F. and Kumarev, V.
Process for preparing polynucleotides on tightly packed bed Patent: US 5869643, A 12,09-FEB-1999;
Unclassified.

1 (bases 1 to 18)

De Rose, R., Douce, R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AR034899
Sequence 18 from patent US
AR034899
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Pred. No. 2e+0
0; Mismatches
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Pred. No. 2e+02;
  Duval,M.,
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2e+02;
    Job, C.
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AR101834
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AR097579/c
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                                                                                                                                                                               Unclassified.

1 (bases 1 to 18)

Manly,S.P., Kozlowski,M.R. and Neve,R.L.
Cloning and expression of .beta.APP-C100

Patent: US 6083713-A 25 04-JUL-2000;
                                                                                                                                                                                                                                                              Unknown
                                                                                                                                                                                                                                                                            Unknown.
                                                                                                                                                                                                                                                                                                                         AR101834
Sequence 25 from patent US
AR101834
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Patent: US 6071745-A 9 06-JUN-2000;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 18)
Lin,C.-I.Patsy., Wallace,R.Bruce., Cossman,J. and French,C.
Method and formulation for lyophilizing cultured human cells to
preserve RNA and DNA contained in cells for use in molecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 9
AR097579
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unclassified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                h 1.1%; Score 14.8; DB 1;
Similarity 88.9%; Pred. No. 2e+02;
16; Conservative 0; Mismatches 2;
                                                                           Similarity
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/mol_type="unassigned DNA"
                                                                                                                     /organism="unknown"
/mol_type="unassigned
                                                                                                                                                                  Location/Qualifiers
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88.9%; Pred. No. 2e+1
vative 0; Mismatches
                                                                         1.1%; Score 14.8; DB 1; 88.9%; Pred. No. 2e+02;
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2e+02;
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RESULT 132
BD222596/c
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AR106909/c
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AR106506
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Best Local Similarity 88.9%;
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Best Local
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AR106506
                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 18)
Cowsert,LM., Bennett,C.Frank. and O'Malley,B.W.
Antisense modulation of SRA expression
Patent: US 6107092-A 70 22-AUG-2000;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AR106909
Sequence 70 from patent US
AR106909
produced therefrom
Patent: JP 2002522447-A 14 23-JUL-2002;
                       synthetic construct synthetic construct artificial sequences. artificial sequences. 1 (bases 1 to 18) 1 (bases 1 to 18) Manoharan, M., Cook, P.D., Prakash, T.P. and Kawasaki, A.M. Aminooxy-modified nucleoside compound and oligomer compound
                                                                                                      produced therefrom.
BD222596
BD222596.1 GI:33032366
JP 2002522447-A/14.
                                                                                                                                                           BD222596 18
Aminooxy-modified nucleoside
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unknown.
Unclassified
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Similarity 88.9%;
16; Conservative
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Patent: US 6107060-A 30 22-AUG-2000;
Location/Qualifiers
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Keeling, P. and Guan, H.
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                                                                                                                                                                                                                                                                                                                                                  /mol_type="unassigned
                                                                                                                                                                                                                                                                                                                                                                 organism="unknown"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              organism="unknown"
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                                                                                                                                                                                                                                                                                               Score 14.8; DB 1;
Pred. No. 2e+02;
0; Mismatches 2;
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Pred. No. 2e+02;
0; Mismatches 2;
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S 6107092.
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                        1520 AAAAAAAAAAAGTAAAA 1537
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Kenichi,H., Hiroshi,Y. and Masahide,N.

Method for labeling oligonucleotide and utilization thereof

Patent: JP 199075880-A 2 23-MAR-1999;

CHEMO SERO THERAPEUT RES INST

OS Unidentified
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                                                                                                                                                                                                                      PI KENICHI HANAKI,HIROSHI YOSHIKURA,MASAHIDE NOZAKI PC
C12N15/09,C12Q1/68,G01N33/58,C12N15/00
CC Strandedness: Single;
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Method for labeling
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Similarity 88.9%;
                                                               n 1.1%;
Similarity 88.9%;
 AAAAAAAAAAAAAAAA 18
                                                                                                                                                                                                                                                                            Unidentified
JP 199075880-A/2
23-MAR-1999
10-JUL-1998 JP 1998195719
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07-AUG-1998 US 09/130973
MUTHIAH MANOHARAN,PHILIP DAN COOK,THAZHA P PRAKASH,ANDREW M
                                                    Conservative
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C12N15/00
                                                                                                                                                                                                           Topology: Linear;
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                                                                                                   /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                                                                    /organism='Unidentified'.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                     GI:13025387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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oligonucleotide
                                                Score 14.8; DB 1;
Pred. No. 2e+02;
0; Mismatches 2;
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Pred. No. 2e+02;
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RESULT 134 E28536/c

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DEFINITION
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                                                      RESULT 136
AR196702/c
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179509/c
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AR196702
Sequence
AR196702
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 16 from patent US 179509
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                                                                                                                                                                                                                                                                                                                                                                                                                 Kato, K
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                      1167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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                                                                                                                                                                                                                                                                                             /mol_type="unassigned DNA"
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                                                                                                                                                                                                                                                                                                                 organism="unknown"
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Pred. No. 2e+0
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Pred. No. 2e+02;
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                    6350934.
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2e+02;
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AR196704/c
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AR215435/c
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                                                                                                                                                                                                                                                                                             AR215435
Sequence 9
AR215435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
                                                        1 (bases 1 to 18)
Lin,C.-I.P., Wallace,R.B., Cossman,J. and French,C.
Method and formulation for lyophilizing cultured human cells to
preserve RNA and DNA contained in cells for use in molecular
biology experiments
Diology experiments
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 18)
Zwick, M.G., Edington, B.E., McSwiggen, J.A., Merlo, P.Ann. Owens.,
Guo, L., Skokut, T.A., Young, S.A., Folkerts, O. and Merlo, D.J.
Nucleic acid encoding delta-9 desaturase
Patent: US 6350934-A 1169 26-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1169 from patent AR196704 AR196704.1 GI:20246141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Zwick, M.G., Edington, B.E., McSwiggen, J.A., Merlo, P.Ann. Owens.,
Guo, L., Skokut, T.A., Young, S.A., Folkerts, O. and Merlo, D.J.
Nucleic acid encoding delta-9 desaturase
Patent: US 6350934-A 1167 26-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unknown
                                                                                                                                                                                                Unclassified
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Similarity 88.9%;
16; Conservative
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Similarity 88.9%; Pred. No. 2e
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/mol_type="unassigned DNA"
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                  organism="unknown"
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AR473365/c
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AR412363/c
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AR222464 AR222464.1 GI:23329995
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Manoharan, M., Cook, P.D., Prakash, T.P. and Kawasaki, A.M. Aminooxy-modified nucleosidic compounds and oligomeric prepared therefrom Patent: US 6639062-A 14 28-OCT-2003;
                                                                                                                                                                                                                                                                                                                          AR412363
Sequence 14 from patent US
AR412363
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1 (Dases 1 to 18)

Kurz,M., Lohse,P. and Wagner,R.

Reptide acceptor ligation methods

Patent: US 6429300-A 24 06-AUG-2002;

Location/Qualifiers
                                                                                                                                                                                                                                                                                Unknown.
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/mol_type="genomic DNA"
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                     from patent US
  GI:42708816
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0; Mismatches
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Pred. No. 2e+02;
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AR487020/c
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Sequence 7
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Sequence 6
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Thirstrup,K., Warthoe,P. and Pettersson,N.B.
Process for amplifying and labeling single s
ligated adaptor mediated amplification
Patent: US 6706476-A 6 16-MAR-2004;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 (bases 1 to 18)
Lin,C.-I.P., Wallace,R.B., Cossman,J. and French,
Method and formulation for lyophilizing cultured
preserve RNA and DNA contained in cells for use i
biology experiments
Patent: US 6686460-A 9.03-FEB-2004;
                                     1 (bases 1 to 18)
Thirstrup, K., Warthoe, P. and Pettersson, N.B
Process for amplifying and labeling single
ligated adaptor mediated amplification
Patent: US 6706476-A 7 16-MAR-2004;
Location/Qualifiers
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/organism="unknown"
/mol_type="genomic |
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Method for isolating anionic organic substances from
systems using cationic polymer nanoparticles
Patent: WO 9910527-A 4 04-MAR-1999;
SUEDDEUTSCHE KALKSTICKSTOFF (DE); BAYER ERNST (DE)
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Sequence 4 from Patent WO9910527.
AXOO4875
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Sequence 8 from Patent WO9910527.
AX004879
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16; Conserv
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Method for isolating anionic organic substances fr
systems using cationic polymer nanoparticles
Patent: WO 9910527-A 8 04-MAR-1999;
SUEDDEUTSCHE KALKSTICKSTOFF (DE); BAYER ERNST (DE)
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/db_xref="taxon:32630"
                                                                                                                    /organism="synthetic construct"
/mal_type="unassigned RNA"
/db xref="taxon:35230"
/noTe="2' methyl-modified oligonucleotide"
                                                                                                                                                                             ocation/Qualifiers
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                                                          Score 14.8; DB Pred. No. 2e+02;
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AX008118/c
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AX008117
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AX008118
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1520 AAAAAAAAAAAGTAAAA 1537
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                                                                                                                                                                                          and its analogues
Patent: WO 9967378-A 3 29-DEC-1999;
DAMHA MASSAD JOSE (CA); PARNIAK MICHAEL A (CA); WILDS CHRISTOPHER
DAMHA MASSAD JOSE (CA); PARNION DOMINIQUE (CA); NORONHA ANNE M (CA)
(CA); UNIV MCGILL (CA); ARION DOMINIQUE (CA);
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DAMHA MASSAD JOSE (CA); PARNIAK MICHAEL
(CA); UNIV_ (CA); ARION DOMINIQUE
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llarity 88.9%;
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/mol_type="unassigned DNA"
/db xref="taxon:32630"
/noTe="Use as an oligomer"
                                                                                                                /organism="synthetic construct"
/mol_type="unassigned RNA"
/db_xref="taxon:32630"
                                                                                                /note="Use as an oligomer"
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Pred. No. 2e+02;
0; Mismatches 2;
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AX028843
Sequence 27 from Patent WO9732023
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Patent: WO 9967378-A 8 29-DEC-1999;
DAMHA MASSAD JOSE (CA); PARNIAK MICHAEL
(CA); UNIV_MCGILL (CA); ARION DOMINIQUE
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Sequence 8 from Patent WO9967378.
AX008123
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Sequence 7 from Patent WO9967378.
AX008122
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Patent: WO 9967378-A 7 29-DEC-1999;
DAMHA MASSAD JOSE (CA); PARNIAK MICHAEL A (CA); WILDS CHRISTOPHER (CA); UNIV MCGILL (CA); ARION DOMINIQUE (CA); NORONHA ANNE M (CA);
BORKOW GADI (IL)
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mal_type="unassigned DNA"
/mb.trype="taxon:32630"
/noce="Use as an oligomer"
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                                                                                                                                                                1.1%; Score 14.8; DB 1; 88.9%; Pred. No. 2e+02;
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KEYWORDS
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AX047273/c
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DEFINITION
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AX047271
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Matches 16
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Sequence 23
AX047273
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Gumbiowski, K. and Zulauf, M.
High density labeling of dna with modified on nucleotides and dna polymerases used Patent: WO 0068422-A 21 16-NOV-2000;
Roche Diagnostics GmbH (DE)
Location/Qualifiers
synthetic construct artificial sequences
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Sequence 21 from Patent WO0068422.
AX047271
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FLORIGENE LIMITED (AU) ; BRUGLIERA FILIPPA
ALBERT (AU) ; MICHAEL MICHAEL ZENON (AU)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Brugliera, F., Holton, T.A. and Michael, M.Z.
Genetic sequences encoding flavonoid pathway enzymes and uses
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AX028843.1 GI:10189946
                                            synthetic construct
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Similarity 88.9%;
16; Conservative
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Similarity 88.9%;
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/mol_type="unassigned DNA"
/mol tyse="claxon:32630"
/moEe="Oligonuclectide"
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/mol_type="unassigned DNA"
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/db_xref="taxon:32630"
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Pred. No. 2e+02;
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High density labeling of dna with modified or chromophore carrying nucleotides and dna polymerases used patent: WO 0068422-A 23 16-NOV-2000;
Roche Diagnostics GmbH (DE)
                                                                                                                                                                                                           AX104747
Sequence 939 from Patent
AX104747
AX104747.1 GI:13920944
                                                                         Krieg,A.M., Schetter,C. and Vollmer,J.C. Immunostimulatory nucleic acids Patent: WO 0122972-A 939 05-APR-2001; UNIVERSITY OF IOWA RESEARCH FOUNDATION (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Krieg,A.M., Schetter,C. and Vollmer,J.C.
Immunostimulatory nucleic acids
Patent: WO 0122972-A 913 05-APR-2001;
UNIVERSITY OF IOWA RESEARCH FOUNDATION (US); Coley Pharmaceutical
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Sequence 10 from Patent WO0123564.
AXI05651
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Sequence 10 from Patent WO0123419.
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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Sequence 913 from Patent
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AX547774.1 GI:25812918
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Sequence 837 from Patent
AX355809
AX355809.1 GI:18620477
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UNIVERSITY OF IOWA RESEARCH FOUNDATION
Location/Qualifiers
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/db_xref="taxon:32630"
/note="Synthetic oligonucleotide~phosphorothioate
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AX547800/c
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Sequence 1 from Patent WO03064441.
AX814716
                                                                                                     Damha,M.J. and Parniak,M.A.
Oligonucleotides comprising alternating segments and uses thereof Patent: WO 03064441-A 1 07-AUG-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Inhibition of angiogenesis by nucleic acids Patent: WO 02053141-A 939 11-JUL-2002; Coley Pharmaceutical Group, Inc. (US) Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   synthetic construct artificial sequences.
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Sequence 939 from Patent
AX547800
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16; Conservative
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                                                                                             UNIVERSITY (CA)
/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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Pred. No. 2e+02;
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AX814723/c
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Sequence 8 from Patent WO03064441.
AX814723
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 l Similarity
16; Conserv
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synthetic construct
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ilarity 88.9%;
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                                                                                                                                                                  UNIVERSITY (CA)
                                                 /note="Residues 1-3, 7-9, 2'-0-methyl-D-uridine"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                  organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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           8; DB 1;
2e+02;
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2e+02;
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AX814725/c
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 RESULT 166
BD085545/c
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AX814736
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AX814725
AX814725.1
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                                                                                                                                                                                                                                                                                                      synthetic construct
synthetic construct
artificial sequences.
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Sequence 21 from Patent WO03064441.
AX814736
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artificial sequences.
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Similarity 88.9%;
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                                                                                                                                                                             /organism="synthetic construct"
/mol_type="unassigned RNA"
/db_xref="taxon:32630"
                                                                                                                                                                 /note="Target RNA oligonucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="Residues 1-6 and 13-18 are 2'-O-methyl-D-uridine"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Pred. No. 2e
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A68209
AR048767
Sequence 1 from
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Collingwood, S.P., Moser, H.E., Altmann, K. and INTERNEDIATES FOR OLIGONUCLEOTIDE SYNTHESIS Patent: WO 9747636-A 4 18-DEC-1997;
                                                                                                                                                                                                                                                                                                              unidentified unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    l Similarity
16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 18)
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JP 2001333800-A/2.
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                                                                                                                                                                                                                                                                                                  unclassified
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4 from Patent WO9747636.
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/mol_type="genomic RNA"
/db_xref="taxon:9606"
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                                                                                                                                                                     /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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88.9%;
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Pred. No. 1.9e+02;
0; Mismatches 2;
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AR111946/c
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AR111371/c
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AR111371
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Leclerc,G. and Martel,R.
Rediolabeled DNA oligonucleotide and method
Patent: US 5821354-A 1 13-OCT-1998;
Location/Qualifiers
                                                                    1 (bases 1 to 19)
Cook, P.Dan., Mancharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 20 03-CCT-2000;
Location/Qualifiers
                                                                                                                                          Unknown.
                                                                                                                                                                              AR111946 1
Sequence 20 from patent US
AR111946
AR111946.1 GI:12828794
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Leeds, J.M. and Cummins, L.L.
Fluorescence based nuclease assay
Patent: US 6127124-A 1 03-OCT-2000;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unknown.
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                                 /organism="unknown"
/mol_type="unassigned DNA"
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02;
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AR111949/c
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AR111948/c
LOCUS
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AR111947/c
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Sequence 21 from patent US 6127533.
AR111947.1 GI:12828795
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            Unknown
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 22 03-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                     Unknown
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 21 03-OCT-2000;
Location/Qualifiers
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Pred. No. 1.9e+02;
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Sequence 24
AR111950
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                                                                                                                                                        1 (bases 1 to 19)
Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 25 03-OCT-2000;
                                                                                                                                                                                                                                                         Sequence 25 from patent US
ARI11951
ARI11951.1 GI:12828799
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 19)
Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 24 03-OCT-2000;
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 23 03-OCT-2000;
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                                                                      1.1%;
Similarity 88.9%;
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/mol_type="unassigned
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                                                                                                                      organism="unknown"
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                                                                                                           _type="unassigned
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AR111957/c
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SOURCE
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ACCESSION
VERSION
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AR111953/c
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AR111952/c
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Sequence 27
AR111953
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1 (bases 1 to 19)

Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.

2'-O-aminooxy-modified oligonucleotides

Patent: US 6127533-A 31 03-OCT-2000;

Location/Qualifiers

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Sequence 31 from patent US 6127533.
AR111957
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 27 03-OCT-2000;
Location/Qualifiers
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.

2'-O-aminooxy-modified oligonucleotides

Patent: US 6127533-A 26 03-OCT-2000;

Location/Qualifiers
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Sequence 26 from patent US
AR111952
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Similarity 88.9%;
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nilarity 88.9%;
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/mol_type="unassigned
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/mol_type="unassigned DNA"
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Pred. No. 1.9e+02;
0; Mismatches 2;
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Pred. No. 1.9e+02;
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Mismatches 2;
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KEYWORDS
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   RESULT 181
AR111970/c
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DEFINITION
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AR111959/c
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AR111970
Sequence 44
AR111970
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AR111959
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Cook, P. Dan., Mancharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 34 03-OCT-2000;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                 Sequence 34 from patent AR111960
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 19)
Cook, P. Dan., Mancharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 33 03-OCT-2000;
Location/Qualifiers
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/mol_type="unassigned
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/mol_type="unassigned
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Pred. No. 1.9e
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Query Match
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Sequence 20
AR124843
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16; Conserv
 l Similarity
16; Conserv
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Unknown.
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1 (bases 1 to 19)
Cook, P.Dan., Manoharan, M. and Kawasaki, A. Mamoru.
2'-O-aminooxy-modified oligonucleotides
Patent: US 6127533-A 44 03-OCT-2000;
                                                                                             Unclassified.

1 (bases 1 to 19)

Manoharan, M., Cook, P.Dan., Prakash, T.P. and Kaw Manoharan, M., Cook, P.Dan., Prakash, T.P. and Kaw Aminooxy-modified oligonucleotides and methods Patent: US 6172209-A 21.09-JAN-2001;
                                                                                                                                                                                                         Sequence 21
AR124844
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Manoharan,M., Cook,P.Dan., Prakash,T.P. and Kawasaki,A.M.
Aminooxy-modified oligonuclectides and methods for making
Patent: US 6172209-A 20 09-JAN-2001;
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                                              /organism="unknown"
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 Score 14.8; D
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                       Length 19;
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AR124845
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Manoharan, M., Cook, P.Dan., Prakash, T.P. and Kawasaki, A.M.
Aminooxy-modified oligonucleotides and methods for making
Patent: US 6172209-A 23 09-JAN-2001;
Location/Qualifiers
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Manoharan, M., Cook
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Sequence 25
AR124848
AR124848.1
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Manoharan, M., Cook, P.Dan., Prakash, T.P. and Kawasaki, A.M.
Aminooxy-modified oligonucleotides and methods for making
Patent: US 6172209-A 26 09-JAN-2001;
Location/Qualifiers
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Sequence 26 from patent US
AR124849
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Manoharan,M., Cook,P.Dan., Prakash,T.P. and Kawasaki,A.M.
Aminooxy-modified oligonucleotides and methods for making
Patent: US 6172209-A 25 09-JAN-2001;
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id Similarity 88.9%;
16; Conservation
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AR124856
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Sequence 27
AR124850
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Manoharan, M. (Cook, P. Dan., Prakash, T. P. Aminooxy-modified oligonucleotides and patent: US 6172209-A 27 09-JAN-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 19)
Manoharan, M., Cook, P. Dan., Prakash, T. P.
Aminooxy-modified oligonucleotides and
Patent: US 6172209-A 31 09-JAN-2001;
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                               1 (bases 1 to 19)
Manoharan, M., Cook, P.Dan., Prakash, T.P.
Aminooxy-modified oligonuclectides and π
Patent: US 6172209-A 33 09-JAN-2001;
                                                                                         Unknown.
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Pred. No. 1.9e+02;
0; Mismatches 2
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Pred. No. 1.9e+02
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                                                  RESULT 194
AR135291/c
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AR124867/c
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AR124857/c
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Best Local Similarity 88.9%;
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 AR135291 1
Sequence 20 from patent US
AR135291
AR135291.1 GI:14124196
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Manoharan, M., Cook, P. Dan., Prakash, T. P. and Kawasaki, A. M. Aminooxy-modified oligonuclectides and methods for making Patent: US 6172209-A 44 09-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                      AR124867
Sequence 44 from patent US
AR124867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unclassified.

1 (bases 1 to 19)

Manoharan, M., Cook, P. Dan., Prakash, T. P. and Kawasaki, A. M. Aminoxy-modified oligonucleotides and methods for making Patent: US 6172209-A 34 09-JAN-2001;
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Sequence 34 from patent US
AR124857
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/mol_type="unassigned DNA"
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02;
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AR135293
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AR135292
                                                                                                  1 (bases 1 to 19)
Cook, P.Dan., Manoharan, M. and Kawasaki, A. Mamoru.
Aminooxy-modified oligonucleotide synthetic intermediates
Patent: US 6194598-A 22 27-FEB-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru. Aminooxy-modified oligonucleotide synthetic intermediates Patent: US 6194598-A 21 27-FEB-2001;
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Aminooxy-modified oligonuclectide synthetic intermediates
Patent: US 6194598-A 20 27-FEB-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unclassified.

1 (bases 1 to 19)
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ilarity 88.9%;
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                                                             /organism="unknown"
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/mol_type="unassigned
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             1.1%; Score 14.8; DB 1
88.9%; Pred. No. 1.9e+02
vative 0; Mismatches
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AR135295/c
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Sequence 23
AR135294
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Sequence
AR135296
Unclassified.
1 (bases 1 to 19)
Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
                                                                                                                                                                                                                                                                                                           1 (bases 1 to 19)
Cook, P.Dan., Manoharan, M. and Kawasaki, A.Mamoru.
Aminooxy-modified oligonucleotide synthetic intermediates
Patent: US 6194598-A 24 27-FEB-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                       Unknown
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Cook, P. Dan., Mancharan, M. and Kawasaki, A. Mamoru.
Aminooxy-modified oligonuclectide synthetic intermediates
Patent: US 6194598-A 23 27-PEB-2001;
                                                                          AR135296.1
                                                                                                                                                                                                                                                                                                                                                                                                   Unknown.
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llarity 88.9%;
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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                    1520 АААААААААААСТААА 1537
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Cook, P.Dan., Mancharan, M. and Kawasaki, A. Mamoru.
Aminooxy-modified oligonuclectide synthetic inte
Patent: US 6194598-A 27 27-FEB-2001;
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Location/Qualifiers
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                                                                                       /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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RESULT 202

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AR135305/c
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1 (cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
Aminooxy-modified oligonucleotide synthetic intermediates
Patent: US 6194598-A 31 27-FEB-2001;
Location/Qualifiers
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Sequence 31
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Cook, P.Dan., Manoharan, M. and Kawasaki, A. Mamoru.
Aminooxy-modified oligonucleotide synthetic inte
Patent: US 6194598-A 34 27-FEB-2001;
Location/Qualifiers
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Sequence 34 from patent US
AR135305
AR135305.1 GI:14124210
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Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru.
Aminooxy-modified oligonucleotide synthetic intermediates
Patent: US 6194598-A 33 27-FEB-2001;
Location/Qualifiers
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Sequence 33 from patent US
AR135304
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llarity 88.9%;
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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AR153863/c
LOCUS
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AR141898/c
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AR153863
AR153863.1 (
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AR141898
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AR135315
AR135315.1
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Manoharan, M., Kawasaki, A.M., Cook, P.Dan., Fraser, A.S.
Prakash, T.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cook, P. Dan., Manoharan, M. and Kawasaki, A. Mamoru. Aminooxy-modified oligonucleotide synthetic inte Patent: US 6194598-A 44 27-FEB-2001;
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Patent: US 6147200-A 4 14-NOV-2000;
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Similarity 88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="unknown"
/mol_type="unassione
                                                                                                                                                                   /mol_type="unassigned
                                                                                                                                                                                                      Location/Qualifiers
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                                                                                                                                                                                 organism="unknown"
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Pred. No. 1.9e+02;
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1.9e+02;
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BD196900/c
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AR164173/c
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JOURNAL
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Best Local Similarity
Matches 16; Conserv
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Sequence 6
AR164173
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PN J
PD 1
PF 2
PR 2
                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eut
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Ho
1 (bases 1 to 19)
Cohen,D., Blumenfeld,M., Chumakov,I. and Bougueleret,L.
Prostatic cancer gene
Patent: JP 2002516657-A 489 11-JUN-2002;
                                                                                                                                                                           BD196900.1 GI:33006670
JP 2002516657-A/489.
Homo Bapiens (human)
                                                                                                                                                                                                                                  BD196900
Prostatic cancer gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 19)
Manoharan,M., Mohan,V. and Boswell,H.
RNA targeted 2'-modified oligonucleotides that are conformationally
preorganized
Patent: US 6271358-A 6 07-AUG-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diagnostics Patent: US 6238624-A 16 29-MAY-2001;
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Heller, M.J., Tu, E., Evans, G.A. and Sosnowski, R.G.
Methods for transport in molecular biological analysis
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Homo sapiens (human)

DP 2002516657-A/489

11-JUN-2002

12-DEC-1998 JP 2000525562

22-DEC-1997 US 08/996306,09-SEP-1998
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Pred. No. 1.9e+02;
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Pred. No. 1.
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Best Local
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PN JP 2002516657-A/500

PD 11-JUN-2002

PF 22-DEC-1998 JP 2000525562

PR 22-DEC-1997 US 08/996306,09-SEP-1998 US 60/099658 PI

DANIEL COHEN, MARTA BLUMENFELD, ILYA CHUMAKOV, LYDIE BOUGUELERET PC

C12N15/09,C12N15/09,A01K67/027,C07K14/47,C07K16/18,C12N1/15, PC

C12N1/19.
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PC C12N5/00,C12N15/00
PC C12N5/00,C12N15/00
PC potential microsequencing oligo for 4-4-187.mis2 FH Key Location/Qualifiers
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PC C12N5/00,C12N15/00
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Cohen, D., Blumenfeld, M., Chumakov, I. and
Prostatic cancer gene
Patent: JP 2002516657-A 500 11-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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                                                                                                                            /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Location/Qualifiers
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                                                              Score 14.8; DB 1;
Pred. No. 1.9e+02;
0; Mismatches 2;
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Pred. No. 1.9e+02;
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RESULT 211 BD274438/c

LOCUS DEFINITION

BD274438 19 l Oligonucleotides having A-DNA

A form

DNA linear PAT 17-JUL-2 and B-DNA form confirmational

PAT 17-JUL-2003

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REFERENCE
AUTHORS
TITLE
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KEYWORDS
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BD274439/c
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  1520 АААЛАЛАЛАЛАЛАГТАЛАЛ 1537
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PATENTS PHARMACEUTICALS INC

OS Artificial Sequence
PN JP 2002543215-A/15

PD 17-DEC-2002

PF 03-MAY-1909 US 009/303

PF 03-MAY-1909 US 09/303

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Oligonucleotides having A-DNA
geometry.
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synthetic construct
artificial sequences.
1 (bases 1 to 19)
Manoharan, M. and Mohan, V.
Oligonucleotides having A-DNA form and B-DNA form
                                                                                                                                                                                                                                                                                                                                                                        synthetic construct
synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 19)
Manoharan,M. and Mohan,V.
Oligonucleotides having A-DNA form and B-DNA form confirmational
                                                                                                                                                                                                                                                                                                                                                              geometry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BD274439.1 GI:33084207 JP 2002543215-A/16.
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JP 2002543215-A/15.
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C12N15/00
Oligonucleotide
2' - O-MOE linkage
2' - O-MOE linkage
2' - O-MOE linkage
                                                                                                                                                                                                                                                       nt: JP 2002543215-A 16 17-DEC-2002;
PHARMACEUTICALS INC
Artificial Sequence
JP 2002543215-A/16
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03-MAY-2000 JP 2000615638
03-MAY-1999 US 09/303586
MUTHIAH MANOHARAN, VENKATRAMAN MOHAN
C07H21/02, A61K48/00, A61P35/00, A61P35/02, A61P43/00, C12N15/09,
                                                                                                                              03-MAY-2000 JP 2000615638
03-MAY-1999 US 09/303586
MUTHIAH MANOHARAN, VENKATRAMAN MOHAN
C07H21/02, A61K48/00, A61P35/00, A61P35/02, A61P43/00, C12N15/09,
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larity 88.9%;
Conservative
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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(17) . . (18)
(18) . . (19) .
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Pred. No. 1.9e
0; Mismatches
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and
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B-DNA form confirmational
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       RESULT 214
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VERSION
KEYWORDS
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BD274440/c
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Best Local
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Best Local :
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                     1520 ААЛАЛАЛАЛАЛАГТАЛАЛ 1537
19
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                                               16;
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1 (bases 1 to 19)
Mancharan,M. and Mohan,V.
Oligonucleotides having A
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JP 2002543215-A/17.
synthetic construct
synthetic construct
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Oligonucleotides having A-DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                 Patent: JP 2002543215-A 17 17-DEC-2002;
                                                          Similarity
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 AAAAAAAAAAAAAAAA
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Artificial Sequence
JP 2002543215-A/17
17-DEC-2002
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misc_feature (16)..(17)
misc_feature (17)..(18)
misc_feature (18)..(19).
Location/Qualifiers
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03-MAY-1999 US 09/301586
MUTHIAH MANOHARAN, VENKATRAMAN MOHAN
C07H21/02,A61K48/00,A61P35/00,A61P35/02,A61P43/00,C12N15/09,
C12N15/00
                                               Conservative
                                                                                                                                                                                                                              3' - O-MOE linkage; sub O linkage
3' - O-MOE linkage; sub O linkage
3' - O-MOE linkage; sub O linkage
3' - O-MOE linkage
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                                                                                          /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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/mol_type="genomic DNA"
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                                             Score 14.8; D
Pred. No. 1.9e
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(17) . . (18)
(18) . . (19)
(19) . . (19)
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Pred. No. 1.9e
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.9e+02;
                                                           .9e+02
                                                                      DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          confirmational
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                                               Gaps
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VERSION
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BD274449/c
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VERSION
KEYWORDS
SOURCE
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SOURCE
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DEFINITION
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Best Local (
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OS Artificial Sequence
PN JP 2002543215-A/26
PD 17-DEC-2002
PD 17-DEC-2002
PF 03-MAY-1999 US 09/
PR 03-MAY-1999 US 09/
PI MUTHIAH MANOHARAN, VEN
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                                                                                           JP 2002543215-A/26.

Bynthetic construct
Bynthetic construct
artificial sequences.

1 (bases 1 to 19)

Manoharan, M. and Mohan, V.

Oligonucleotides having A-DNA form and B-DNA form confirmational
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   synthetic construct
synthetic construct
artificial sequences.

1 (bases 1 to 19)
Manoharan,M. and Mohan,V.
Oligonucleotides having A-DNA form and B-DNA
                                                                         geometry Patent: JP 2002543215-A 26 17-DEC-2002;
                                                                                                                                                                                                         BD274449
Oligonucleotides having
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JP 2002543215-A/18.
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Oligonucleotides having A-DNA
                                                                                                                                                                          BD274449.1 GI:33084217
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Similarity 88.9%;
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misc_feature
misc_feature
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03-MAY-2000 JP 2000615638
03-MAY-1999 US 09/303586
MUTHIAH MANOHARAN, VENKATRAMAN MOHAN
C07H21/02,A61K48/00,A61P35/00,A61P35/02,A61P43/00,C12N15/09,
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PHARMACEUTICALS INC
Artificial Sequence
JP 2002543215-A/18
 17-DEC-2002
03-MAY-2000 JP 2000615638
03-MAY-1999 US 09/303586
03-MAY-1999 US VENKATRAMAN MOHAN
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misc_feature
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2'- O-MOE; sub O linkage
2'- O-MOE; sub O linkage
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                                                                                                                                                                                                                                                                                                                                                             /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Pred. No. 1.9e+02;
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                                                                                                                                                                                                         DNA linear
and B-DNA form
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AUTHORS
TITLE
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VERSION
KEYWORDS
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AUTHORS
TITLE
                                                                                                                                                    RESULT 217
AR205799/c
LOCUS
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ORGANISM
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VERSION
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AR205798/c
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Best Local
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1520 АААААААААААСТААА 1537
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Sequence 15
AR205798
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Sequence 16
AR205799
geometry
Patent: US 6369209-A 16 09-APR-2002;
Location/Qualifiers
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Patent: US 6369209-A 15 09-APR-2002;
Location/Qualifiers
                                                1 (bases 1 to 19)
Manoharan,M. and Mohan,V.
Oligonucleotides having A-DNA form and B-DNA form conformational
                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 19)
Mancharan, M. and Mohan, V.
Oligonucleotides having A-DNA form and
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                                                                                               Unclassified.
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2'-modified T linkage
2'-modified T linkage
2'-modified T linkage
2'-modified T linkage
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/mol_type="genomic DNA"
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                                                                                                                                                                                                                                                                                                                                                                                  /mol_type="unassigned
                                                                                                                                                                                                                                                                                                                                                                                                /organism="unknown"
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                                                                                                                                                       GI:21503473
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(17) . . (18)
(18) . . . (19)
(19) . . . (19)
                                                                                                                                                                                                                                                                                                                          0; Mismatches
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Pred. No. 1.9e+02;
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Pred. No. 1.9e+02
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6369209.
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RESULT 220
AR205809/c
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AR205800/c
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AR205801/c
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Matches 16
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Patent: L
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Sequence 18 from patent US
AR205801
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Manoharan,M. and Mohan,V.
Oligonucleotides having A-DNA form and B-DNA form conformational
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AR205800
  AR205809
                                                                                                                                                                                                   Unclassified.
1 (bases 1 to 19)
Manoharan,M. and Mohan,V.
Oligonucleotides having A-DNA form and B-DNA
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                                                                                                                                     /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Pred. No. 1.9e+02;
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Pred. No. 1.9e.
0; Mismatches
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AR213490/c
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1520 ААААААААААААСТАААА 1537
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Sequence 1 from I
AR213490
AR213490.1 GI:23
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Sequence 2
AR213491
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Patent: US 6403779-A 1 11-JUN-2002;
Location/Qualifiers
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Manoharan,M. and Mohan,V.
Oligonucleotides having A-DNA form and B-DNA form conformational
                        Regioselective synthesis of 2'-O-modified nucleosides
Patent: US 6403779-A 2 11-JUN-2002;
Location/Qualifiers
                                                                        1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
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Kawasaki,A.M., Fraser,A.S., Manoharan,M., Cook,P.D. and
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                                                                                                                         Unclassified
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Similarity 88.9%; Pred. No. 1.9e+02;
16; Conservative 0; Mismatches 2;
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Similarity 88.9%;
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/mol_type="genomic DNA"
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/mol_type="unassigned DNA"
/organism="unknown"
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Pred. No. 1.9e+02;
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TITLE
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AR213494/c
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SOURCE
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AR213493/c
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AR213492/c
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AR213494
Sequence 5 from
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Sequence 4 from patent US
AR213493
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                                                                                                                                                                                                                        Unclassified.

(bases 1 to 19)

Kawasaki,A.M., Fraser,A.S., Manoharan,M., Cook,P.D. and Prakash,T.P.
                                                                                                                                                                                                                                                                     Unknown
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1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
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llarity 88.9%;
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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DEFINITION
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AR213496/c
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AR213496
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                                              Regioselective synthesis of 2'-O-modified Patent: US 6403779-A 7 11-JUN-2002;
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Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
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Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
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Kawasaki, A.M., Fraser, A.S., Manoharan, M., Cook, P.D. and Prakash, T.P.
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Similarity 88.9%; Pred. No. 1.99
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AR213497/c
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AR213502/c
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Sequence 14 from patent US 6403779.
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Sequence 12 from patent US
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1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
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Sequence 8 from patent US 6403779.
AR213497 GI:23310728
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Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
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Pred. No. 1.9e+02;
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Patent: US 6403779-A 15 11-JUN-2002;
Location/Qualifiers
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AR213512
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Kawasaki A.M., Fraser, A.S., Manoharan, M.,
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Kawasaki A.M., Fraser, A.S., Manoharan, M.,
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Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
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Sequence
AR222465
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Kurz,M., Lohse,P. and Wagner,R.

Peptide acceptor ligation methods

Patent: US 6429300-A 25 06-AUG-2002;

Location/Qualifiers
                      AR295557
Sequence 7292 from patent
AR295557
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Sequence 1 from patent US
AR237463
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           AR295557.1
                                                                                                                                                                                                                                        Process for the synthesis of oligomeric Patent: US 6465628-A 1 15-OCT-2002; Location/Qualifiers
                                                                                                                                                                                                                                                                            Unclassified.
1 (bases 1 to 19)
Ravikumar, V.T., Manoharan, M., Capaldi, D.C., Krotz, A.,
Guzaev, A.
                                                                                                                                                                                                                                                                                                                             Unknown
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ilarity 88.9%;
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/mol_type="genomic DNA"
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            GI:31682841
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Pred. No. 1.9e+02;
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AR359804/c
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AR321589/c
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Sequence 10
AR321589
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Baxter,A.D., Collingwood,S.P., Douglas,M.E. and Taylor,R.J.
Oligonucleotide analogues
Patent: US 6562960-A 10 13-MAY-2003;
Location/Qualifiers
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Cohen, D., Chumakov, I. and Blumenfeld, M.
Biallelic markers for use in constructing a high density
disequilibrium map of the human genome
Patent: US 6537751-A 7292 25-MAR-2003;
                                                                                           1 (bases 1 to 19)
Manoharan,M. Cook,P.D., Prakash,T.P. and Mohan,V.
Guanidinium functionalized nucleotides and precurs
Patent: US 6593466-A 3 15-JUL-2003;
Location/Qualifiers
                                                                                                                                                                                                                   AR359804
Sequence 3 from patent US
AR359804
             16;
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/mol_type="genomic DNA"
                                                           /mol_type="genomic
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                                                                       organism="unknown"
            1.1%; Score 14.8; DB 1;
88.9%; Pred. No. 1.9e+02;
vative 0; Mismatches 2;
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AR359805/c
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AR367447/c
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AR359806/c
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Sequence 4 :
AR367447
AR367447.1
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Sequence
AR359805
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Mancharan,M., Cook,P.D., Prakash,T.P. and Mohan,V.
Guanidinium functionalized nucleotides and precurs
Patent: US 6593466-A 5 15-JUL-2003;
Location/Qualifiers
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Sequence 5 from patent I
AR359806
AR359806.1 GI:33766604
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Manoharan,M., Cook,P.D., Prakash,T.P. and Mohan,V. Guanidinium functionalized nucleotides and precurs Patent: US 6593466-A 4 15-JUL-2003;
1 (bases 1 to 19)
Collingwood, S.P., Moser, H.E.,
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    Altmann, K.-H.
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    and Douglas, M.E.
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AR399178/c
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Sequence
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AR399177
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Crooke, S.T., Lima, W.F., Wu, H. and Monoharan, M.
Human Rnase H1 and oligonucleotide composition
Patent: US 6617442-A 18 09-SEP-2003;
                                                                                                                                                                                                     Unknown
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Crooke, S.T., Lima, W.F., Wu, H. and Monoharan, M.
Human Rnase H1 and oligonucleotide composition
Patent: US 6617442-A 17 09-SEP-2003;
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                                                                                                                                                                                        Unclassified.
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Similarity 88.9%; Pred. No. 1.9e+02;
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/mol_type="genomic |
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/mol_type="genomic DNA"
                                                                                        /organism="unknown"
/mol_type="genomic |
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                                             Score 14.8; D
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RESULT 243

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AR403602/c
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AR403603/c
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AR403602
AR403602.1 GI:40151188
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Sequence 1 from patent US
AR403601
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Sequence 3 from patent US
AR403603
AR403603.1 GI:40151189
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Regioselective synthesis of 2'-O-modified Patent: US 6624294-A 3 23-SEP-2003; Location/Qualifiers
                                   1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
                                                                                           Unknown.
                                                                                                                                                                                                                                                                                                                                                                           Unclassified.
1 (bases 1 to 19)
Kawasaki, A.M., Fraser, A.S., Manoharan, M., Prakash, T.P.
                                                                                                                                                                                                                                                                                                                                                                                                                         Unknown
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Drawasaki, A.M., Fraser, A.S., Mancharan, M.,
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ilarity 88.9%;
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/mol_type="genomic DNA"
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Pred. No. 1.9e+02;
0; Mismatches 2
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Pred. No. 1.9e+02;
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                                                 Cook, P.D.
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AR403604/c
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AR403605/c
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AR403604
AR403604.1 GI:40151190
                                                                                                                                Regioselective synthesis of 2'-O-modified nucleosides Patent: US 6624294-A 5 23-SEP-2003; Location/Qualifiers
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Sequence 5 from patent US
AR403605
AR403605.1 GI:40151191
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1 (bases 1 to 19)

1 (bases 1 to 19)

Kawasaki, A.M., Fraser, A.S., Manoharan, M., Cook, P.D. ar Prakash, T.P.

Regioselective synthesis of 2'-O-modified nucleosides Patent: US 6624994-A 23-SEP-2003;
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                                                                                                                                                                       Unclassified.
1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
                                                                                                                                                                                                                         Unknown.
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Similarity 88.9%;
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Similarity 88.9%;
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Pred. No. 1.9e+02;
                                                           Score 14.8; DB 1;
Pred. No. 1.9e+02;
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AUTHORS
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AR403608/c
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AR403607/c
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Local Similarity 88.9%;
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Patent: US 6624294-A 6 23-SEP-2003;
Location/Qualifiers
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Sequence 7 from patent US
AR403607
        Regioselective synthesis of 2'-O-modified nucleosides
Patent: US 6624294-A 8 23-SEP-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                               Regioselective synthesis of 2'-O-modified Patent: US 6624294-A 7 23-SEP-2003; Location/Qualifiers
                                            Unclassified.

1 (bases 1 to 19)

Kawasaki, A.M., Fraser, A.S., Manoharan, M., Cook, P.D. and Prakash, T.P.
                                                                                          Unknown
                                                                                                       Unknown.
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Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
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Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
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                                                                                                                                                                                                                                                                                                             /organism="unknown"
/mol_type="genomic DNA"
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from patent US 6624294.
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88.9%;
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Pred. No. 1.9e+02;
0; Mismatches 2
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Pred. No. 1.9e+02;
0; Mismatches 2
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3 6624294.
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          RESULT 253
AR403614/c
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AR403613/c
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Sequence
AR403613
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Sequence
AR403612
                                                                                                        16;
                                                                                                                                                                                                                     1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 19)
Kawasaki, A.M., Fraser, A.S., Manoharan, M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16;
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 AR403614
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Similarity 88.9%;
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/mol_type="genomic
                                                                                                                                                                                     Location/Qualifiers
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/mol_type="genomic |
                                                                                                                                                   /organism="unknown"
/mol_type="genomic DNA"
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Pred. No. 1.9e+02;
0; Mismatches 2;
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Pred. No. 1.9e+02;
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Length 19;

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Gaps

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Cook, P.D. and

Cook, P.D. and

Length

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AR412338/c
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AR403623/c
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Sequence
AR403623
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                                  1 (bases 1 to 19)
Cook, P.D., Manoharan,M., Maier,M. and An,H.
C3'-methylene hydrogen phosphonate oligomers and
Patent: US 6639061-A 1 28-OCT-2003;
Location/Qualifiers
                                                                                                                                                             Sequence 1 from AR412338
                                                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 19)
Kawasaki,A.M., Fraser,A.S., Manoharan,M.,
Prakash,T.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Regioselective synthesis of 2'-O-modified Patent: US 6624294-A 15 23-SEP-2003;
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1 (bases 1 to 19)

Kawasaki, A.M., Fraser, A.S., Manoharan, M., Prakash, T.P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unknown
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/organism="unknown"
/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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AR451262/c
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Sequence 7
AR432617
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1 (bases 1 to 19)

Manoharan,M., Cook,P.D. and Guinosso Manoharan,M., Cook,P.D. and Guinosso Modified oligonucleotides

Patent: US 6653458-A 6 25-NOV-2003;

Location/Qualifiers
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AR432616
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Manoharan, M., Cook, P.D. and Guinoss
Modified oligonucleotides
Patent: US 6653458-A 7 25-NOV-2003;
                                                                                                                                                                                                                                                               Unclassified.
                                                                                                                                                                                                                                                                         Unknown.
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Similarity 88.9%;
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/mol_type="genomic DNA"
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/mol_type="genomic"
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           GI:42682240
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Pred. No. 1.9e+02;
0; Mismatches 2
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AX059378/c
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Mancharan,M. and Cook,P.D.
2'-O-aminoethyloxyethyl-modified oligonucleotides
Patent: US 6673912-A 5 06-JAN-2004;
Location/Qualifiers
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Sequence 26 from patent US 6673912.
AR451282
                                                                                                 Plant chromosome compositions and methods Patent: WO 0055325-A 111 21-SEP-2000; The University of Chicago (US)
                                                                                                                                                                   Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Sequence 111 from Patent WO0055325.
AX059378 GI:12311483
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Manoharan, M. and Cook, P.D.

2'-O-aminoethyloxyethyl-modified oligonucleotides

Patent: US 6673912-A 26 06-JAN-2004;
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/mol_type="genomic DNA"
                            /organism="Arabidopsis thaliana"
/mol_type="unassigned DNA"
/db_xref="taxon:3702"
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/mol_type="genomic DNA"
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Ribozyme therapy for the treatment of proliferative skin and
                                                                                                                                                              Patent: WO 0160856-A 52 23-AUG-2001;
UNIVERSITE CATHOLIQUE DE LOUVAIN (BE)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                /organism="synthetic construct"
/mol_type="unasigned DNA"
/mol_type="unasigned DNA"
/db xrefs"ttaxon:32630"
/note="oligonucleotide"
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Pred. No. 1.9e+02
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PF 01-DEC-
PR 05-DEC-
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BD087505
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synthetic construct
artificial sequences.
1 (bases 1 to 19)
1 (bases, R.G., Butler, W.F., Tu, E., Nerenberg, M.I., Heller, M.J.
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Patent: WO 0202810-A 33 10-JAN-2002;
Clondiag Chip Technologies GmbH (DE)
Location/Qualifiers
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Method for qualitative and/or quantitative detecting of molecular
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AX349249
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JP 2001525193-A/16.
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17-2001525193-A/16
11-DEC-2001
01-DEC-1998 JP 2000524303
05-DEC-1997 US 08/986065
05-DEC-1997 US 08/986065
05-DEC-1997 US 08/986065
                                                                                                                                                                              C12Q1/68, C12N15/09, C12N15/00
Description of Artificial Sequence:
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llarity 88.9%;
Conservative
                                                                                                                                                                                                                   MICHAEL J HELLER, CARL F EDMAN
                                                    /organism='Artificial Sequence'
/organism="synthetic construct"
/mol_type="genomic DNA"
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/mol_type="unassigned DNA"
/db xrefe"taxon:32630"
/noTe="Oligonukleotidsonde"
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from Patent WO0202810.
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Pred. No. 1.9e+02;
0; Mismatches 2;
                                                                                                         Location/Qualifiers
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AR064875/c
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AR030917/c
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                                                                                                                                                                                                                                                                     Unknown
                                                                                                                                                                                                           Process and device for assaying a hapten Patent: US 5849480-A 5 15-DEC-1998;
                                                                                                                                                                                                                                1 (bases 1 to 20)
Cros, P., Kurfurst, R., Battail, N. and Piga, N.
                                                                                                                                                                                                                                                                                                                 Sequence 5
AR064875
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AR030917
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genetic sequences encoding flavonoid pathway enzymes and uses
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Holton, T. Albert.,
Lester, D. Ruth.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unclassified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unknown.
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                                                                       AAAAAAAAAAAAAAA 3
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20 bp
83 from patent US 5968524.
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                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="unassigned DNA"
                                                                                                                                                             /mol_type="unassigned DNA"
                                                                                                                                                                         organism="unknown"
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88.9%; Pred. No. 1.9e+02;
vative 0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                       GI:5995091
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                                                                                                                   <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
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Pred. No. 1.8e+02;
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Pred. No. 1.
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Mismatches 2
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S 5861487.
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AR087520/c
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AR085926
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Sequence 1
AR087520
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Sequence 83
AR085926
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Patent: US 5968524-A 83 19-OCT-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                    16; Conserv
                                     i (bases 1 to 20)
Pitsch,S., Weiss,P.A. and Jenny,L.
Ribonucleoside-derivative and method for preparing the
Patent: US 596084-A 1 16-NOV-1999;
Location/Qualifiers
                                                                                                                                                                                                             μ
                                                                                                 Unknown.
                                                                                                                                                                                                                                                                                                                        Patent: US 5985287-A 83 16-NOV-1999;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 20)
Tan, P., Skinner, M. and
Compounds and methods
                                                                                                                                                                                                                                                                                                                                                                                            Unknown
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Watson, J.D. and Tan, P.
Methods and compounds
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AR080000.1
                                                                                        Unclassified.
                                                                                                            Unknown.
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                                                                                                                                                                                                                                                                                        /mol_type="unassigned
          /mol_type="unassigned
                   organism="unknown"
                                                                                                                                                                                                                                                                                                    organism="unknown"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mol_type="unassigned DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                organism="unknown"
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                                                                                                                                GI:10014283
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Pred. No. 1.8e+02;
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                              RESULT 272
AR121540
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AR118970/c
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1520 AAAAAAAAAAAGTAAAA 1537
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 (bases 1 to 20)
Tan,P., Hiyama,J., Visser,E., Skinner,M.,
Mycobacterium vaccae antigens
Patent: US 6001361-A 83 14-DEC-1999;
Location/Qualifiers
                                                                                                                                                                                                                  1 (bases 1 to 20)
Uchida, K., Uchida, T., Tanaka, Y., Matsuda, Y.
Antieense nucleic acid compound targeted to
Patent: US 6150092-A 96 21-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                1 AAAAAAAAAAAAAAAA
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AR121540.1
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Similarity 88.98;
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                                                                                                                                                                      /organism≃"unknown"
/mol_type="unassigned
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/mol_type="unassigned
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AR123335
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AR121692
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Best Local Similarity 88.9%;
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Sequence 83
AR121692
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McKay, R., Borchers, A.H. and Baker, B.F.

Antisense modulation of peroxisome proliferator-activated receptor

gamma expression

Patent: US 6159734-A 76 12-DEC-2000;

Location/Qualifiers
 16; Conserv
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Bruice, T.C. and Dev, A.P.
Deoxynucleic alkyl thioures compounds and
Patent: US 6169176-A 1 02-OAN-2001;
Location/Qualifiers
                                                                                                                                                                                         AR123335
Sequence 1 from patent US
AR123335
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Location/Qualifiers
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tive 0; Mismatches 2;
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AR139962/c
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Unclassified.
1 (bases 1 to 20)
Zsebo, K.M., Bosselman, R.A.,
                                          Unknown.
                                                                                            AR140279
Sequence 32 from patent
AR140279
                                                                                                                                                                                                                                                                                                                                                                                                                       Unknown
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Zsebo,K.M., Bosselman,R.A., Suggs,S.V.
Zsebo,K.M.n stem cell factor
DNA encoding stem cell factor
Patent: US 6207417-A 32 27-MAR-2001;
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AR139960
                                                      Unknown.
                                                                               AR140279.1
                                                                                                                                                                                                                                                                                                                                                     DNA encoding stem cell factor Patent: US 6207417-A 34 27-MAR-2001;
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Zsebo, K.M., Bosselman, R.A., Suggs, S.V.
                                                                                                                                                                                                                                                                                                                                                                                                        Unclassified.
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nilarity 88.9%;
Conservative
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                                                                                 GI:14482775
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S 6207454.
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6207417.
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  Suggs, S.V. and Martin, F.H.
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AR140557/c
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Sequence 32 from patent US
AR140557
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                                                  16;
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Spebo, K.M., Bosselman, R.A., Suggs, S.V. and Martin, F.H.

Method for enhancing the effciency of gene transfer with stem factor (SCF) polypeptide

Patent: US 6207454-A 34 27-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 34 from patent US AR140281
                                                                                                                                                     1 (bases 1 to 20)
Zsebo, K.M., Bosselman, R.A., Suggs, S.V.
Stem cell factor and compositions
Patent: US 6207802-A 32 27-MAR-2001;
                                                                                                                                                                                                                   Unknown
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Similarity 88.9%;
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                                                                                                 /organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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                                              Score 14.8; DB 1;
Pred. No. 1.8e+02;
Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
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AR154115/c
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AR141070/c
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AR140559/c
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Sequence 1
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Sequence
AR154115
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AR140559
Huang, Z. and Szostak, J.W.
Simple and efficient method to label and modify 3'-termini of RNA using DNA polymerase and a synthetic template with defined overhar nucleotides
                                                                                                                                                                                                                                               16;
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Manoharan, M. and Maier, M.A.
Compounds, processes and intermediates
backbone oligomeric compounds
Patent: US 6207819-A 1 27-MAR-2001;
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Zsebo, K.M., Bosselman, R.A., Suggs, S.V.
Stem cell factor and compositions
Patent: US 6207802-A 34 27-MAR-2001;
Location/Qualifiers
                                                                     Unknown
                                                                               Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                            Unknown
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1 (bases 1 to
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/mol_type="unassigned
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Pred. No. 1.8e+02;
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BD218101
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Best Local Similarity 88.9%;
Matches 16; Conservative
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AL PATENT: JP 2002514385-A 26 21-MAY-2002;
GRNESIS RESEARCH AND DEVELOPMENT CORP LTD

OS ARTIficial Sequence
PN JP 2002514385-A/26
PD 21-MAY-2002
PF 23-DEC-1998 JP 2000525553
PR 23-DEC-1997 US 08/997362,23-DEC-1997 US 08/997080 PR
23-DEC-1997 US 08/996624,11-JUN-1998 US 09/95855 PR
23-DEC-1997 US 08/956624,11-JUN-1998 US 09/205426 PI PAUL
17-SEP-1998 US 09/155181,04-DEC-1998 US 09/205426 PI PAUL
17-SEP-1998 US 09/155181,04-DEC-1998 US 09/205426 PI PAUL
17-SEP-1998 US 09/155181,04-DEC-1998 US 09/205426 PI PAUL
18-PRESTICIOE
PI L PRESTICIOE
PC C12N15/09,A61K31/711,A61K39/04,A61K48/00,A61P11/00,A61P11/00,PC A61P17/00,A61P31/00,A61P31/00,A61P37/04,C07K14/35,C07K16/12,PC C07K19/00,
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Blumberg, B.
High through
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Sequence 13
AR164658
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                                                                                                                                                                                                                                                                                    artificial sequences.

1 (bases 1 to 20)

Tan,P., Watson,J., Visser,E.S., Skinner,M.A. and Prestid,R.L.

Compositions derived from mycobacterium vaccae and methods for
                                                                                                                                                                                                                                                                                                                                                           synthetic construct
                                                                                                                                                                                                                                                                                                                                                                                           BD218101.1 GI:33027871 JP 2002514385-A/26.
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Compositions derived from
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     A61P17/00,
A61P17/06,A61P31/00,A61P31/06,A61P37/04,C07K14/35,C07K16/12,
C07K19/00,
                                                       L PRESTIDGE
C12N15/09, A61K31/711, A61K39/04, A61K48/00, A61P11/00, A61P11/06,
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Location/Qualifiers
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/mol_type="unassigned
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mycobacterium vaccae and methods
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BD234126
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Sequence 40
CQ759610
CQ759610.1
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PC (C12N15/09,C12R1:32),C12N15/00,C12N5/00,(C12N15/00,C12R1:3
PT Key Location/Qualifiers
FT source /organism='Artificial Sequence'.
                                                                                                                                         Н
                                                                                                                                                                                              l Similarity 88.9
16; Conservative
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synthetic construct
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Protein skeleton of antibody
synthetic construct
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1 (bases 1 to 20)
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Similarity 88.9%;
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C12N15/00
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C12N15/09,C07K1/04,C07K14/78,C07K16/46,C07K17/00,C07K19/00,
                                                                                                                                                                                                                                                                                                                                                                                                                                     Artificial Sequence
JP 2002532072-A/14
02-0CT-2002
09-DEC-1999 JP 2000587187
10-DEC-1998 US 60/111737
                                                                                                                                                                                                                                                                                                                                                   Puromycin linker oligonucleotide
Key Location/Qualifiers
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Location/Qualifiers
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/mol_type="genomic DNA"
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Location/Qualifiers
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/mol_type="genomic DNA
/db_xref="taxon:32630"
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88.9%;
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Pred. No. 1.8e+02;
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Pred. No. 1.8e+02
                                                      20 bp DNA
WO2003106672.
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SOYAKU GIJUTSU KENKYUSHO:KK
OS NONE
OC Artificial sequences.
PN JP 1997052898-A/10
PD 25-FEB-1997
PF 09-AUG-1995 JP 1995224608
PI MIZUGUCHI MASATSUGU, KURO
KOYANAGI YOSHIO,
PI YAMAMOTO NAOKI
PC COTHR1/04/A61K31/70;
CC strandedness: Single;
CC topology: Linear;
CC hypothetical: No;
CC antisense: Yes;
FH Key Location,
FT Source 1..20
FT Source 1..20
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Mizuguchi, M., Kurosaki, N., Makino, K., Koyanagi, Y. and Yamamoto, N.
ANTI-HTLY-I ANTI-SENSE OLIGONUCLECTIDE
Patent: JP 1997052898-A 10 25-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E12676.1 GI:325150 JP 1997052898-A/10.
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Patent: WO 2003106672-A 40 24-DEC-2003;
Riken (JP) ; Kabushiki Kaisha Dnaform (JP)
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MIZUGUCHI MASATSUGU, KUROSAKI NAOKO,
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/noTe="tag8"
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/db_xref="taxon:32644"
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Pred. No. 1.8e+02;
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147310/c
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Sequence 11 from patent US 5639870.
147310
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Sequence 16 from patent US
I36180
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Holton,T.A., Cornish,E.C., Kovacic,F., Tanaka,Y. and Lester,D. Genetic sequences encoding flavonoid pathway enzymes and uses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 from pate
128309
128309.1 GI:1819085
                                                 Holton,T.Albert., Cornish,E.Cecily. and Tanaka,Y. Genetic sequences encoding flavonoid pathway enzy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unknown
                                                                 1 (bases 1 to 20)
Holton, T. Albert.,
                                                                                                        Unknown
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Heller, M.J. and Tu, E.
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                          Patent: US 5639870-A 11 17-JUN-1997;
                                         therefor
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20 from patent US 5569832.
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Location/Qualifiers
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/mol_type="unassigned
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/mol_type="unassigned
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AR231312/c
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AR222466
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Sequence
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AR231312
Sequence
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1 (bases 1 to 20)
Kurz,M., Lohse,P. and Wagner,R.
Peptide acceptor ligation methods
Patent: US 6429300-A 26 06-AUG-2002;
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1 (bases 1 to 20)

Tan,P., Visser,E., Prestidge,R. and Watson,J.D.

Compounds and methods for treatment and diagnosis of mycobacterial
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Sequence 83 from patent US
AR213738
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llarity 88.9%;
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49 from patent US
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/mol_type="unassigned DNA"
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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Pred. No. 1.8e+02;
0; Mismatches 2;
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                                                                                                      Score 14.8; DB 1;
Pred. No. 1.8e+02;
0; Mismatches 2
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S 6406704.
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6451968.
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6429300.
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AR274394
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I (bases 1 to 20)

Egholm,M., Nielsen,P., Buchardt,O., I Coull,J.M., Kiely,J. and Griffith,M. Peptide nucleic acids

Patent: US 6451968-A 49 17-SEP-2002;

Location/Qualifiers
                                                                   1 (bases 1 to 20)
Mirkin, C.A., Letsinger, R.L., Mucic, R.C., Elghanian, R. and Taton, T.A.
Nanoparticles having oligonucleotides att
                                                                                                                                                                                                        Sequence 55
AR274394
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Compounds, processes and intermediates backbone oligometic compounds patent: US 6462184-A 1 08-OCT-2002; Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unknown.
                                                                                                                                                  Unknown.
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Manoharan, M. and Maier, M.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unknown.
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AR236083
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                             Patent: US 6506564-A 55 14-JAN-2003;
Location/Qualifiers
                                                                                                                                  Unclassified.
                                                                                                                                                               Unknown.
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/mol_type="genomic DNA"
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/mol_type="genomic
/organism="unknown"
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84.2%; Pred. No. 1.8e+02;
tive 0; Mismatches
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Pred. No. 1.8e+02
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                                                                          attached thereto and uses
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                                                                                                      Storhoff, J.J.,
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AR365970
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l Similarity 88.9%;
16; Conservation
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AR365970 20 bp
Sequence 83 from patent US 6328978.
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                                                                                                                                                                                                                             Patent: US 6582921-A 55 24-JUN-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                       (bases 1 to 20)
Mirkin,C.A., Letsinger,R.L., Mucic,R.C., Storhoff,J.J.,
Elghanian,R. and Taton,T.A.
Nanoparticles having oligonucleotides attached thereto and uses
thereof
                                                                                                                                                                                                                                                                                                                                       Unknown
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Sequence 55 from patent US
AR344936
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Manoharan, M., Lonnberg, H., Salo, H. and Virta, P. Aminooxy functionalized oligomers
Patent: US 6576752-A 10 10-JUN-2003;
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/mol_type="genomic DNA"
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/mol_type="genomic |
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88.9%; Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
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AR429653
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                                      1 (bases 1 to 20) Mirkin,C.A., Letsinger,R.L., Mucic,R.C., Elghanian,R. and Taton,T.A. Nanoparticles having oligonucleotides att therefor
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AR429653
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Mirkin,C.A., Letsinger,R.L., Mucic,R.C., Storhoff,J.J.,
Elghanian,R. and Taton,T.A.
Nanoparticles having oligonucleotides attached thereto and uses
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AR382312
AR382312.1 GI:40090724
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Watson,J.D., Tan,P.L.J. and Prestidge,R.
Methods for the treatment of immunologically-mediated skin
                                                                                                                         Unknown
                                                                                                                                     Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent: US 6328978-A 83 11-DEC-2001;
Location/Qualifiers
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AR365970.1
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Location/Qualifiers
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  Patent: US 6645721-A 55 11-NOV-2003;
Location/Qualifiers
1. .20
                                                                                                         Unclassified.
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Similarity 88.9%;
16; Conservative
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/mol_type="genomic DNA"
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Pred. No. 1.8e+02;
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                                                         attached thereto and uses
                                                                                  Storhoff, J.J.,
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PAT 18-DEC-2003

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PAT 18-DEC-2003

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RESULT 302
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Best Local Similarity
Matches 16; Conserv
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Sequence 55
AR451990
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Sequence 55 from patent US
AR447441
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                                                                                                                                 Patent: US 6677122-A 55 13-JAN-2004;
Location/Qualifiers
                                                                                                                                                                    1 (bases 1 to 20)
Mirkin,C.A., Letsinger,R.L., Mucic,R.C., Storhoff,J.J.,
Elghanian,R. and Taton,T.A.
Nanoparticles having oligonucleotides attached thereto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unclassified.

1 (bases 1 to 20)
Mirkin,C.A., Letsinger,R.L., Mucic,R.C., Storhoff,J.J.,
Elghanian,R. and Taton,T.A.
Nanoparticles having oligonucleotides attached thereto
                                                                                                                                                                                                                                    Unknown.
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ilarity 88.9%;
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ilarity 88.9%;
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
                                                                                             /organism="unknown"
/mol_type="genomic |
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Pred. No. 1.8e-
0; Mismatches
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                                               Score 14.8; DB 1;
Pred. No. 1.8e+02;
0; Mismatches 2
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RESULT 303

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Sequence 7 :
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Sequence 55
AR454776
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Sequence 55
AR489044
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1 Similarity 88.9%;
16; Conservative
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16; Conserv
1 (Dases 1 to 20)
1 (Dases 1 to 20)
Mirkin,C.A., Letsinger,R.L., Mucic,R.C.,
Elghanian,R. and Taton,T.A.
Nanoparticles having oligonucleotides att
                                                               Unknown
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Nelson, W.G., Lin, X., Tchou, J.C. and Bakker, J.
Methods of diagnosing and treating hepatic ce
                                                                                                                                                                                                                                                                                                                                                                                                                Unknown.
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Mirkin, C.A., Letsinger, R.L., Mucic, R.C.,
Elghanian, R. and Taton, T.A.
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                                                  Unclassified
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                                                                                                 AR489044.1
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ilarity 88.9%;
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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Pred. No. 1.8e+02;
0; Mismatches 2
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      attached thereto
                             Storhoff, J.J.,
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FEATURES
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AR494728
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AR494116
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Best Local Similarity
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                      1520 ААААААААААААСТАААА 1537
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1 (bases 1 to 20)
Mirkin, C.A., Letsinger, R.L., Mucic, R.C., Storhoff, J.J., Elghanian, R. and Taton, T.A.
Nanoparticles having oligonucleotides attached thereto
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                                                                                                                              Patent: US 6720411-A 55 13-APR-2004;
Location/Qualifiers
                                                                                                                                                                1 (bases 1 to 20)
Mirkin,C.A., Letsinger,R.L., Mucic,R.C., Storhoff,J.J.,
Elghanian,R. and Taton,T.A.
Nanoparticles having oligonucleotides attached thereto
                                                                                                                                                                                                                          Unknown
                                                                                                                                                                                                                                                                          Sequence 55 from patent US AR494728
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Location/Qualifiers
                                                                                           /mol_type="genomic DNA"
                                                                                                       organism="unknown"
                                                                                                                                                                                                                                                                                                                                                                                                                                                     /mol_type="genomic DNA"
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Pred. No. 1.8e+02;
0; Mismatches 2;
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                                            Score 14.8; DB 1;
Pred. No. 1.8e+02;
0; Mismatches 2
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AUTHORS
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AX045779/c
                                   RESULT 310
AX045787/c
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AX004876/c
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AX045787
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16; Conserv
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Sequence 9 from Patent W00067023.
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Method for isolating anionic organic substances
systems using cationic polymer nanoparticles
Patent: WO 9910527-A 5 04-MAR-1999;
SUEDDEUTSCHE KALKSTICKSTOFF (DE); BAYER ERNST (ED)
Location/Qualifiers
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Sequence 5 from Patent WO9910527.
AX004876
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                               /note="modified with digoxigenin"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Pred. No. 1.8e+02;
0; Mismatches 2;
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Murinae gen. sp.
Murinae gen. sp.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                           Sequence 241 from Patent AX056597
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Screening for immunostimulatory dna functional modifyers
Patent: WO 0067023-A 20 09-NOV-2000;
CPG Immunopharmaceuticals GmbH (DE); UNIVERSITY OF IOWA RESEARCH
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Sequence 20 from Patent W00067023.
AX045790
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Screening for immunostimulatory dna functional modifyers
Patent: WO 0067023-A 17 09-NOV-2000;
CPG Immunopharmaceuticals GmbH (DE); UNIVERSITY OF IOWA RESEARCH
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llarity 88.9%;
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|mol_type="unassigned DNA"
|db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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RESULT 313
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AX104364/c
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Patent: WO 0073469-A 241 07-DEC-2000;
Sugen, Inc. (US)
                                                                                                                                                                                                                    Sequence 556 from Patent AX104364
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                                                                                Krieg, A.M., Schetter, C. and Vollmer, J.C. Immunostimulatory nucleic acids Patent: WO 0122972-A 556 05-APR-2001; UNIVERSITY OF IOWA RESEARCH FOUNDATION (
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
/db_xref="taxon:39108"
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Pred. No. 1.8e+02;
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Matches 16; Conserv
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16; Conserv
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                                                        1.1%;
l Similarity 88.9%;
16; Conservative
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Nanoparticles having oligonucleotides attached thereto and uses
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Sequence 55 from Patent WO0151665.
AX196224
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Immunostimulatory nucleic acids
Patent: WO 0122972-A 560 05-APR-2001;
UNIVERSITY OF IOWA RESEARCH FOUNDATION (US); Coley Pharmaceutical
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                                                                                                                                                                           Nanosphere,
                                                                                                                                                                                       Patent: WO 0151665-A 55 19-JUL-2001;
                                                                                                                                                                                                                                                                        synthetic construct
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                                  AAAAAAAAAAAGTAAAA 1537
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            AAAAAAAAAAAAAAA 18
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                                                                                                     /organism="synthetic construct"
/mol type="unassigned DNA"
/db xref="taxon.32630"
/noTe="random synthetic sequence"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                                                                            e, Inc. (US)
Location/Qualifiers
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Pred. No. 1.8e
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Pred. No. 1.8e+02;
0; Mismatches 2;
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Pred. No. 1.8e+02;
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AX196239
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KEYWORDS
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RESULT 319
AX355810/c
LOCUS
DEFINITION
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AX354974
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Query Match
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Matches 16;
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Sequence 838 from Patent WO0197843.
AX355810
AX355810.1 GI:18620478
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Sequence 2 from Patent W00197843.
AX354974
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Nanoparticles having oligonucleotides attached thereto
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AX196239
AX196239.1 GI:15386442
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UNIVERSITY OF IOWA RESEARCH FOUNDATION
Location/Qualifiers
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1.1%; Score 14.8; DB 1; Length
Similarity 88.9%; Pred. No. 1.8e+02;
16; Conservative 0; Mismatches 2; Indels
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/noTe="Synthetic oligonucleotide-phosphodiester backbone"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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Pred. No. 1.8e+02;
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AX440125
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Best Local Similarity
Matches 16; Conserv
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Sequence 55
AX440125
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16; Conserv
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UNIVERSITY OF IOWA RESEARCH FOUNDATION (US)
Location/Qualifiers
Nanoparticles
therefor
                           Mirkin, C.A., Letsinger, R.L., Mucic, R.C., Storhoff, J.J., Elghanian, R., Taton, T.A., Park, S.J. and Li, Z.
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UNIVERSITY OF IOWA RESEARCH FOUNDATION
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Sequence 839 from Patent
AX355811
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Methods for enhancing antibody-induced cell lysis
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llarity 88.9%;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:32630"
/note="Synthetic oligonucleotide~phosphodiester backbone"
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Pred. No. 1.8e.
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Pred. No. 1.8e+02;
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AX465311
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Sequence 55 from Patent WO0218643.
AX465311
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                                                                                                                       Mirkin,C.A., Letsinger,R.L., Mucic,R.C., Storhoff,J.J., Elghanian,R., Taton,T.A., Garimella,V., Li,Z. and Park, Nanoparticles having oligonucleotides attached thereto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nanosphere, Inc.
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Nanoparticles having oligonucleotides attached thereto and uses
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                                                           Nanosphere, Inc. (US)
Location/Qualifiers
                                                                                            Patent: WO 0218643-A 55 07-MAR-2002;
                                                                                                                                                                                artificial sequences.
                                                                                                                                                                                                 synthetic
                                                                                                                                                                                                             synthetic construct
                                                                                                                                                                                                                                            AX465311.1 GI:21899674
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/db_xref="caxon:32630"
/noTe="random synthetic sequence"
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/db_xref="taxon:32630"
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Pred. No. 1.8e+02;
0; Mismatches 2
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                                                                                                                                                                              Maruyama, T., Ishiguro, T. and Taya, T. Oligonucleotide and method for detecting verotoxin Patent: EP 1236806-A 8 04-SEP-2002;
                                                                                                                                                                                                                                        synthetic construct synthetic construct
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                                                                                    /note="Oligonucleotide capable of binding specifically VT2 RNA"
                                                                                                        /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
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                                                Score 14.8; DB 1;
Pred. No. 1.8e+02;
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Li,Z. and Park,S.J.
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AX547417/c
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Inhibition of angiogenesis by nucleic a
Patent: WO 02053141-A 556 11-JUL-2002;
Coley Pharmaceutical Group, Inc. (US)
Location/Qualifiers
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Pred. No. 1.8e+02;
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TITLE
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Best Local Similarity 88.9%;
Matches 16; Conservative
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                                                                                                                                                               AX556139 20 bp
Sequence 70 from Patent WO0246472.
AX556139
                                       Mirkin, C.A., Letsinger, R.L., Mucic, R.C., Storhoff, J.J., Elghanian, R., Taton, T.A., Garimella, V., Li, Z. and Park, S.J. Nanoparticles having oligonucleotides attached thereto and
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Patent: WO 0246472-A 70 13-JUN-2002; Nanosphere, Inc. (US)
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/note="Synthetic Sequence"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                      /note="random synthetic sequence"
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Pred. No. 1.8e+02;
0; Mismatches 2
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AX664307
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AX664308/c
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Patent: WO 0246398-A 5 13-JUN-2002;
The University of Houston System (US)
Location/Qualifiers
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Patent: WO 0246398-A 6 13-JUN-2002;
The University of Houston System (University of Houston System)
                                                                                                                                                                                   Willson,R.C. and Murphy,J.C. Nucleic acid separation using immobilized
                                                                                                                                                                                                                                                                                                            Sequence 6 from Patent WO0246398
AX664308
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                                                              /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Pred. No. 1.8e+02;
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Pred. No. 1.8e+02;
    Score 14.8; DB 1;
Pred. No. 1.8e+02;
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TITLE
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Matches 16; Conserv
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                                                                                                                                                                                                                                    Kirtsen,N.V., Hyldig-Nielsen,J.J. and Williams,B.F.
Methods, kits and compositions pertaining to the suppression of detectable probe binding to randomly distributed repeat sequences in genomic nucleic acid
Patent: WO 03027328-A 14 03-APR-2003;
Boston Probes, Inc. (US); DakoCytomation Denmark A/S (DK)
Location/Qualifiers
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Sequence 14 from Patent WO03027328.
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/mol_type="unassigned DNA"
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                                                                                                                                                                                                           organism="synthetic construct"
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Query Match
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                                                                                                                                                                                                                                                       Patent: JP 2001503969-A 26 27-MAR-2001;
GENESIS RESEARCH & DEVELOPMENT CO LTD
OS Unidentified
PN UP 2001503969-A/26
PD 27-MAR-2001
PF 28-AUG-1997 ID 1000FT157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 AAAAAAAAAAAAAAAA 18
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Methods, kits and compositions pertaining to the suppression of detectable probe binding to randomly distributed repeat sequences in genomic nucleic acid patent: WO 03027328-A 26 03-APR-2003;
Boston Probes, Inc. (US), DakoCytomation Denmark A/S (DK)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     synthetic construct
synthetic construct
artificial sequences.
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Sequence 26 from Patent WO03027328.
AX741052
AX741052.1 GI:30523913
                                                                                                                                                                                                                                                                                                                                                            Tan,P., Hiyama,J., Visser,E.S., Skinner,M.A.,
Prestidge,R.L.
Compounds and methods for treatment and diagno
                                                                                                                                                                                                                                                                                                                                                                                                                                                     JP 2001503969-A/26. unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Compounds and methods for infections.
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ROSS L PRESTIDGE
A61KX39/04, A61K35/74, C07K14/35, C12N15/63
Strandedness: Single;
                                                                                                                                                                                                                                                     Unidentified
JP 2001503969-A/26
27-MAR-2001
28-AUG-1997 JP 1998511516
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                                                                                                                      source
                                                                                                                                               opology: Linear
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/db_xref="taxon:32630"
/note="Description of Combined DNA/RNA Molecule:Synthetic Oligomer Sequence-Synthetic Probe Sequence"
                                                                                    /organism='Unidentified'
                         /organism="unidentified'
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Pred. No. 1.8e+02;
0; Mismatches 2;
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 Score 14.8;
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Scott, L.M.

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PAT 31-JAN-2002 Mycobacterial

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BD107450/c
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BD080522/c
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TITLE
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JOURNAL
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Best Local Similarity 88.9%;
Matches 16; Conservative
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DN BD107450.1 GI:23202268

DN BD107450.1 GI:23202268

SP 2002034599-A/9.

SUIT 2002034599-A/9.

NISM Synthetic construct

Artificial sequences.

1 (bases 1 to 20)

ORS Segawa,M. Takarada,H., Aono,T. and Yoshiga,S.

E Method of detecting single base polymorphism

NALL PROVIDED TO LTD

OS Artificial Sequence

PN JP 2002034599-A/9

PD 05-FEB-2002

PF 26-JUL-2000 JP 2000225354

PI MASAYA SEGAWA,HIROSHI TAKARADA,TOSHIYA AONO,SATOKO YOSHIGA PC

C1201/68,C12N15/09,C12N15/09

CC Description of Artificial Sequence:primer

FH Key

FT source 1.20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        artificial sequences.

E 1 (bases 1 to 20)

S Pitsch, S., Weiss, P.A. and Jenny, L.

Pitsch, S., Weiss, P.A. and Jenny, L.

Ribonucleoside-derivative and method for preparing the sam.

Patent: JP 2001515087-A 1 18-SEP-2001;

STEFAN PITSCH, PATRICK A WEISS, LUZI JENNY

OS Artificial Sequence
PN JP 2001515087-A/1

PD 18-SEP-2001

PD 18-SEP-2001

PR 18-AUG-1998 JP 2000509723

PR 18-AUG-1998 JP 200509723

PR 18-AUG-1997 CH 1931/97

PI STEFAN PITSCH, PATRICK A WEISS, LUZI JENNY

PC C07H19/06, C07F7/18, C07H19/16, C07H21/02, C07H23/00 CC
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D 18-SEP-2001
P 17-AUG-1998 JP 2000509723
PR 18-AUG-1997 CH 1931/97
PI STEFAN PITSCH, PATRICK A WEISS, LUZI JENNY
PC C07H19/06,C07F7/18,C07H19/16,C07H21/02,C07H23/00 CC
Description of Artificial Sequence:synthetic polynucleot:
Location/Qualifiers
Location/20
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JP 2001515087-A/1.
synthetic construct
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Ribonucleoside-derivative
BD080522
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/mol_type="genomic RNA"
/db_xref="taxon:32630"
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Pred. No. 1.8e
O; Mismatches
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AR080294/c
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BD174543/c
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Sequence 13
AR080294
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synthetic construct
artificial sequences.
1 (bases 1 to 20)
1 (bases 1 to 20)
Maruyama, T., Ishiguro, T. and Taya, T.
Oligonucleotide for detecting Vero toxin and detection method
Patent: JP 2002253257-A 8 10-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PI TAKAHIRO MARUYAMA, TAKAHIKO ISHIGURO, TOSHITAKA TAYA PC C12N15/09, C12Q1/68, G01N33/53, G01N33/566, C12N15/00 CC Oligonucleotide capable of binding specifically to VT2 RNA
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Oligonucleotide for detecting
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                                                1 (bases 1 to 21)
Watson, M.A. and Fleming, T.P.
Mammaglobin, a mammary-specific breast
Patent: US 5968754-A 13 19-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BD174543.1 GI:29120233 JP 2002253257-A/8.
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Similarity 88.9%;
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JP 2002253257-A/8
10-SEP-2002
02-MAR-2001 JP 2001058143
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA
/db_xref="taxon:32630"
       /organism="unknown"
                                    Location/Qualifiers
                                                                                                                                                                         GI:10007029
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Pred. No. 1
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Pred. No. 1.8e+02;
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Fing Vero toxin and detection method.
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AR084524/c
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1.1 Similarity 88.9%;
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AR093143 2
Sequence 12 from patent US
AR093143
AR093143.1 GI:10019895
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Sequence 10 from patent US
AR084521
                                                                                                                                                                                                                                  1 (bases 1 to 21)
Matson, R.S., Coassin, P.J., Rampal, J.B.
Oligonucleotide repeat arrays
Patent: US 5981185-A 13 09-NOV-1999;
                                                                                                                                                                                                                                                                                                                                          AR084524
Sequence 13 from patent US
AR084524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 (bases 1 to 21)
Matson, R.S., Coassin, P.J., Rampal, J.B.
Oligonucleotide repeat arrays
Patent: US 5981185-A 10 09-NOV-1999;
                                                                                                                                                                                                                                                                                              Unknown.
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Similarity 88.9%;
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                                                                                                                                                                                    organism="unknown"
/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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                                                                                                                                                    Score 14.8; DB 1;
Pred. No. 1.7e+02;
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Pred. No. 1.7e+02;
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Pred. No. 1.7e+02;
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AR095412/c
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AR095412
                                                                 1 (bases 1 to 21)
Brooks-Wilson,A.R., Buckler,A., Cardon,L.,
Miller,A. and North,M.
Asthma related genes
Patent: US 6087485,A 66 11-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 21)
Watson,M.A. and Fleming,T.P.
Method for detecting the presence of breast increase in mammaglobin mRNA expression Patent: US 6004756-A 13 21-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 21)
Bergan,R. and Neckers,L.
Inhibition of protein kinase activity
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AR103542
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Patent: US 5998596-A 12 07-DEC-1999;
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/mol_type="unassigned DNA"
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                         /mol
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                       _type="unassigned
  1.1%;
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AR153849
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TITLE
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Matches 16; Conserv
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Mammaglobin, breast cancer s
BDZ24108
BDZ24108.1 GI:33033878
JP 2002525998-A/10.
Synthetic construct
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Heller,M.J., Tu,E., Evans,G.A. and
Methods for transport in molecular
diagnostics
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AR118155
                                                                                                                                                                                                                                                                                      Patent: US 6238624-A 2 29-MAY-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                     Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2 from patent US AR153849
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Brenner, S.
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Unclassified.
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pred. No. 1.7e+02;
0; Mismatches 2;
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1; Mismatches 3;
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artificial sequences.
E 1 (bases 1 to 21)
E 1 (bases 1 to 21)
S Watson, M.A. and Fleming, T.P.
Mammaglobin, breast cancer secretory protein specific to Patent: JP 2002525098-A 10 13-AUG-2002;
WASHINGTON UNIVERSITY
OS Artificial Sequence
PN JP 2002525098-A/10
PD 13-AUG-2002
PF 29-SEP-1998 US 09/162622
PR 29-SEP-1998 US 09/162622
PR 29-SEP-1998 US 09/162622
PR MARK A WATSON, TIMOTHY P FLEMING
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Sequence 2
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CC Descri
FH Key
FT source
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Sequence 46 from Patent WO2004036221.
CQ846797
CQ846797.1 GI:50895947
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                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                         Patent: WO 2004036221-A 46 29-APR-2004;
Wyeth (US); O'Toole, Margot Mary (US);
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                              O'Toole, M.M. and Liu, W. Compositions and methods for diagnosing and treating autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C12N15/09, C12Q1/68, G01N33/53, G01N33/566, G01N33/577//G01N33/574,
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Similarity 88.9%;
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/mol_type="unassigned RN
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
     GI:2086679
                               from patent US
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Heller, M.J. and Tu, E.
Active programmable electronic devices for molecular biological analysis and diagnostics
Patent: US 5605662-A 2.25-FEB-1997;
                                                                             Massively parallel sequencing of sorted Patent: US 5695934-A 23 09-DEC-1997;
                                                                                                   1 (bases 1 to 21)
Brenner, S.
                                                                                                                                     Unknown.
                                                                                                                                                Unknown.
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Watson, M.A. and Fleming, T.P.
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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Watson,M.A. and Fleming,T.P.
Watson,M.A. and erected mammary-specific breast cancer protein Mammaglobin, a secreted mammary-specific breast cancer protein Patent: US 6677428-A 13 13-JAN-2004;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (bases 1 to 21) / Watson, M.A. and Fleming, T.P. Watson, M.A. and Fleming, T.P. Mammaglobin, a secreted mammary-specific breast cancer protein Patent: US 6566072-A 13 20-WAY-2003; Location/Qualifiers
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88.9%; Pred. No. 1.7e+02;
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Sequence 840 from Patent
AX355812
                                                     Inhibition of angiogenesis by nucleic acids Patent: WO 02053141-A 912 11-JUL-2002; Coley Pharmaceutical Group, Inc. (US)
                                                                                                                                   synthetic construct
synthetic construct
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UNIVERSITY OF IOWA RESEARCH FOUNDATION (US)
Location/Qualifiers
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                                                                                            Bratzler, R.L
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llarity 88.9%;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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                                                                                                                                                                                                                            Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucleic acids Patent: WO 03072818-A 23 04-SEP-2003; Degussa Bioactives GmbH (DE)
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Sequenz:Capture-Oligonukleotid"
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/db_xref="taxon:32630"
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Pred. No. 1.7e+02;
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AX825128/c
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Sequence 25 from Patent W003072818.
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AX825127.1 GI:39750856
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Sequence 26 from Patent WO03072818.
AX825128
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                                                                      Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucleic acids Patent: WO 03072818-A 26 04-SEP-2003; Degussa Bioactives GmbH (DE)
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
/organism="synthetic construct"
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/mod_base=OTHER
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|/mol_type="unassigned DNA"
|/db_xref="taxon:32630"
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AX825130/c
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Patent: WO 03072818-A 26 04-SEP-2003;
Degussa Bioactives GmbH (DE)
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Sequence 29 from Patent WO03072818.
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Sequence 28 from Patent W003072818.
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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AX825132/c
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Sequence 30 from Patent W003072818.
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AX825132.1 GI:39750861
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                                                                                                                                                                                                                                                                                                           synthetic construct synthetic construct artificial sequences.
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Location/Qualifiers
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sequenz:Capture-Oligonukleotid"
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Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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/mod_base=OTHER
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                               note="LNA-T (Locked Nucleic Acid)"
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nilarity 88.9%;
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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/mod_base=OTHER
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/mod_base=OTHER
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Sequence 32 from Patent W003072818.
AX825134
AX825134.1 GI:39750863
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Location/Qualifiers
                                                                             Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucleic Patent: WO 03072818-A 37 04-SEP-2003; Degussa Bioactives GmbH (DE)
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              /organism="synthetic construct"
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/db_xref="taxon:32630"
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Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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/mol_type="unassigned |
/db_xref="taxon:32630"
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/note="Beschreibung
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Sequence 38 from Patent WO03072818.
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AX825140.1 GI:39750869
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llarity 88.9%;
Conservative
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Sequenz:Capture-Oligonukleotid"
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/note="LNA-T (Locked Nucleic Acid)"
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                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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mod_base=OTHER
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/mod_base=OTHER
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                                            mod_base=OTHER
                                                                                                                          note="LNA-T (Locked Nucleic Acid)"
mod_base=OTHER
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                                                            note="LNA-T (Locked
                                                                                                   note="LNA-T (Locked Nucleic Acid)"
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; Pred. No. 1.7e+
0; Mismatches
                                                            Nucleic
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1.7e+02;
ches 2;
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AX825142/c
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synthetic construct synthetic construct artificial sequences.
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Sequence 40 from Patent WO03072818.
XX825142
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Method for sorting single-stranded nucle
Patent: WO 03072818-A 39 04-SEP-2003;
Degussa Bioactives GmbH (DE)
Location/Qualifiers
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Sequence 39 from Patent W003072818.
AX825141
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'mod_base=OTHER
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mod_base=OTHER
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Pred. No. 1.7e+02;
0; Mismatches 2
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AX825143/c
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Sequence 41 from Patent W003072818.
AX825143
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Sequenz:Capture-Oligonukleotid"
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|note="Beschreibung der kuenstlichen
|sequenz:Capture-Oligonukleotid"
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Pred. No. 1.7e+02;
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RESULT 375
AX825144/c
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Sequence 42 from Patent W003072818.
AX825144
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nilarity 88.9%;
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/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
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                                                     ; Score 14.8; D;
; Pred. No. 1.7e
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                                                                    1.7e+02;
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RESULT 376

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Sequence 44 from Patent WO03072818
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Sequence 43 from Patent W003072818.
AX825145
                                                                                                         Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucleic acids Patent: WO 03078818-A 44 04-SEP-2003; Degussa Bioactives GmbH (DE)
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/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
/note="Beschreibung der kuenstlichen Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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Sequence 45 from Patent W003072818.
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                /note="LNA-T (Locked Nucleic Acid)"
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Sequence 46 from Patent WO03072818.
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Sequenz:Capture-Oligonukleotid"
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Sequence 48 from Patent
AX825150
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Location/Qualifiers
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/mol_type="unassigned DNA"
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Sequenz:Capture-Oligonukleotid"
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Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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/mod_base=OTHER
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/mod_base=OTHER
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mod_base=OTHER
                'note="LNA-T (Locked
                                                                                                   note="LNA-T (Locked Nucleic Acid)"
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Pred. No. 1.7e+02;
0; Mismatches 2
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RESULT 383
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Sequence 54 from Patent WO03072818.
AX825156
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/mod_base=OTHER
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Sequenz:Capture-Oligonukleotid"
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/db_xref="taxon:32630"
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/mod_base=OTHER
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Pred. No. 1.7e+02;
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Sequence 55 from Patent W003072818.
AX825157
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AX825158
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Sequenz:Capture-Oligonukleotid"
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db_xref="taxon:32630"
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/mod_base=OTHER
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/mol_type="unassigned DNA"
/db_xref="taxoon:32630"
/noTe="Beschreibung der kuenstl
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                                 /note="Beschreibung der kuenstlichen
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mod_base=OTHER
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Best Local Similarity 88.9%;
Matches 16; Conservative
Query Match
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Sequence 58 from Patent WO03072818.
AX825160
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/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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                                                          'mod_base=OTHER
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                                       note="LNA-T (Locked Nucleic Acid)"
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Sequence 60 from Patent WO03072818.
AX825162
AX825162.1 GI:39750891
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Sequence 59 from Patent WO03072818.
AX825161
                                                                     synthetic construct synthetic construct
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Boekenkamp,D., Dieck,T.H. and Hoppe,H.U. Method for sorting single-stranded nucleic acids
                                                      artificial sequences
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="LNA-T (Locked Nucleic Acid)"
/mod_base=OTHER
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/mod_base=OTHER
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Sequence 62 from Patent WO03072818.
AX825164
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Bioactives GmbH (DE)
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                                                                                                                                                                                    /organism="synthetic construct"
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/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Sequenz:Capture-Oligonukleotid"
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/mod_base=OTHER
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Location/Qualifiers
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Sequence 63 from Patent
AX825165
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Similarity 88.9%; Pred. No. 1.7e+02;
16; Conservative 0; Mismatches 2
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/mod_base=OTHER
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Sequenz:Capture-Oligonukleotid"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          note="LNA-T (Locked Nucleic Acid)"
                                                                                                                                                        88.98;
                                                                                                                                           0;
                                                                                                                                                                          Score 14.8;
                                                                                                                                           Pred. No. 1.7e+02;
Nismatches
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      DNA
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      linear
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PAT 11-DEC-2003

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Matson, M.A. and Fleming, T.P.

Mammaglobin, a secreted mammary specific brea

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COMMENT

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AUTHORS
TITLE
JOURNAL
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BD080832/c
LOCUS
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KEYWORDS
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ACCESSION
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Best Local S
Matches 16
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16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   synthetic construct synthetic construct
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AX825166
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                                               18-SEP-1998 JP 2000511779
18-SEP-1997 US 08/933149
MARK A WATSON TIMOTHY P FLEMING
C12N15/09,A61K35/26,A61K39/00,A61K39/00,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/395,A61K39/39
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/mol_type="unassigned DNA"
/ppe="taxon:32630"
/note="Beschreibung der kuenstlichen
sequenz:Capture-Oligonukleotid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /mod_base=OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       note="LNA-T (Locked Nucleic Acid)"
|mod_base=OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 note="LNA-T (Locked Nucleic Acid)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           note="LNA-T (Locked Nucleic mod_base=OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          note="LNA-T (Locked Nucleic Acid)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8; DB 1
Pred. No. 1.7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 bp
mammary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             specific breast cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Indels
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ACCESSION
VERSION
KEYWORDS
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                                                                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE
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Best Local
                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AUTHORS
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                            1520 AAAAAAAAAAAAGTAAAA 1537
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OS Artific
PN JP 2001
PD 11-DEC-
PF 01-DEC-
PR 05-DEC-
PI RONALD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BD087491 21 bp DNA linear PAT 27-AUG-2002 Self-assembling microelectronic integration system capable of designating self address, compartment device, mechanism, method and operation for molecular biological analysis and diagnosis.

BD087491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          l Similarity
16; Conserv
۳,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 21)
Sosnowski,R.G., Butler,W.F., Tu,E., Nerenberg,M.I., Heller,M.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SSSEFF
                                                                                                                                                                                                 T S S S S S S E E E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       designating microelectronic integration system capable of designating self address, compartment device, mechanism, method operation for molecular biological analysis and diagnosis Patent: JP 2001525193-A 2 11-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                NERENBERG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JP 2001525193-A/2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BD087491.1 GI:22633101
                                                                         Similarity
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АААААААААААААААА 18
                                                                                                                                                                                                                                                                                                                                                                                                        Artificial Sequence
JP 2001525193-A/2
11-DEC-2001
                                                                                                                                                                                                                                                        provide ribonucleic acid base for reactivity; Poly A sequence
    for reduced
                                                                                                                                                                                                                                                                                                  MICHAEL J HELLER, CARL F EDMAN
C12Q1/68, C12N15/09, C12N15/00
Description of Artificial Sequence: Synthesized with u at 3'
                                                                                                                                                                                                                                                                                                                                                              01-DEC-1998 JP 2000524303
05-DEC-1997 US 08/986065
RONALD G SOSNOWSKI,WILLIAM F BUTLER,EUGENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Topology: Linear;
Topology: Linear;
Mammaglobin, a secreted mammary specific breast
Location/Qualifiers
Key
                                                          Conservative
                                                                                                                                                                                                               source
                                                                                                                                                                                                                             Key
                                                                                                                                                                                                                                           secondary structure
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                                                                                                                                                                              /organism='Artificial
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism='Unidentified'.
Location/Qualifiers
                                                                                                             /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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                                                                        1.1%;
88.9%;
                                                                                                                                                                                                                                                                                       terminus to
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                                                        0; Mismatches
                                                                                                                                                                                                                           Location/Qualifiers
                                                                        Score 14.8; DB 1
Pred. No. 1.7e+02
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Pred. No. 1.7e+02;
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                                                                                       DB 1;
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                                                                                      Length
                                                                                                                                                                                               Sequence'
                                                             Indels
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A79657/c
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Best Local Similarity
Matches 15; Conserv
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Best Local Similarity
Matches 16; Conserv
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Patent: JP 2002500895-A 62 15-JAN-2002;

AXYS PHARMACEUTICALS INC
SUDIDENTIFIED
PN JP 2002500895-A/62
PN JP 2002500895-A/62
PD 15-JAN-2002
PD 15-JAN-2002
PD 15-JAN-1998 JP 2000528715
PI ANGELA R BROOKS WILSON, ALAN BUCKLER, LON
CARDON, ALISOUN H CARBY,
PI MARGARET GALVIN, ANDREW MILLER, MICHAEL NORTH
PI MARGARET GALVIN, ANDREW MILLER, MICHAEL NORTH
PI C1201/68, A01K67/027, C07K14/47, C12N15/09, C12N15/00 CC
Strandedness: Single;
CC Topology: Linear;
CC Topology: Linear;
CC Asthma-associated gene
FH Key
FT source

/organism='Unidentified'.
                            1518 ТТАААААААААААСТААА 1536
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 19
                                                                                                                                                                                 1 (bases 1 to 19)
Emrich, T. and Leying, H.
METHOD OF DETECTING TELOMERASE ACTIVITY
PALENT: WO 9720069-A 6 05-JUN-1997;
BOEHRINGER MANNHEIM GMBH (DE); EMRICH TI
                                                                                                                                                                                                                                                                                                                                  A79657 19 bp
Sequence 6 from Patent WO9720069.
A79657
                                                                                                                                                                                                                                                                            unidentified
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Wilson, A.R.B., Buckler, A.,
Miller, A. and North, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BD129772.1 GI:23224717 JP 2002500895-A/62.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="unidentified"
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/db_xref="taxon:32644"
                                                                                                               /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
                                                                                                                                                                       Location/Qualifiers
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                                                                 1.0%;
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                                                       Score 14.6; DB 1;
Pred. No. 2.2e+02;
2; Mismatches 2
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Pred. No. 1.7e+02;
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AX825107
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AR147331/c
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synthetic construct
artificial sequences.
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AX825107
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Emrich, T., Leying, H., Hinzpeter, M. and Karl, G. Method of detecting telomerase activity
Patent: US 6221594-A 6 24-APR-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                              Bookenkamp, D., Dieck, T.H. and Hoppe, H.U. Method for sorting single-stranded nucle Patent: WO 3072818-A 5 04-SEP-2003; Degussa Bioactives GmbH (DE)
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AR147331
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                                     /note="LNA-T (Locked Nucleic Acid)"
/mod_base=OTHER
                                                                                                                       /mod
                                                                                                                                                                                                                                                                    Sequenz:Capture-Oligonukleotid"
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/db_xref="taxon:32630"
/note="Beschreibung der ku
                                                                            note="LNA-T (Locked
mod_base=OTHER
                                                                                                                                                            mod_base=OTHER
                                                                                                                                                                                                  note="LNA-T (Locked Nucleic Acid)"
mod_base=OTHER
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             note="LNA-T (Locked Nucleic Acid)"
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from Patent WO03072818.
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78.9%; Pred. No. 2
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                                                                                                                  RESULT 398
BD085544/c
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AX825117
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Method of comparison and d

BD085544

BD085544.1 GI:22631154

BD085544.1 GI:22631154
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Sequence 15
AX825117
AX825117.1
Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
1 (bases 1 to 22)
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17; Conserv
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llarity 81.0%;
Conservative
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/mod_base=OTHER
18
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Beschreibung der kuenstlichen
Seguenz:Capture-Oligonukleotid"
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from Patent WO03072818.
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detection
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            Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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                                                                     AR089217
Sequence 13
AR089217
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17; Conserv
                       Unknown
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            Unclassified
                                                            AR089217.1
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                                                                                                                                                                                                                                                /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                              Inc. (US)
                                                                                                                                                                                                                                                                                               Location/Qualifiers
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/mol_type="genomic RNA"
                                                                                                                                                                                                               1.0%;
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                                                            GI:10015974
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Method of comparison and detection of RNA amount and Patent: JP 2001333800-A 1 04-DEC-2001;

UNITECH CO LTD

OS Homo sapiens (human)
PN JP 2001333800-A/1
PD 04-DEC-2001
PF 30-MAY-2000 JP 2000160324
PI KAORI SHIMADA
PC C1201/68,C11N15/09,G01N33/50,C12N15/00
CC Method of comparison and detection of RNA amounced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AX692829 25 bp D
Sequence 5561 from Patent EP1281758.
AX692829
AX692829.1 GI:29415792
                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                     Patent: EP 1281758-A 5561 05-FEB-2003;
                                                                                                                                                                                                                                                                                                                                                                                          Shannon, M., Gu, Y. and Nguyen, C.T. Four human zinc-finger-containing proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens (human)
JP 201333800-A/1
04-DEC-2001
30-MAY-2000 JP 2000160324
KAORI SHIMADA
C12Q1/68,C12N15/09,G01N33/50,C12N15/00
Method of comparison and detection of RNA amount and DNA
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Pred. No. 1.9e+02
0; Mismatches
                                                                                                                                                               Score 14.6; DB 1;
Pred. No. 1.6e+02;
0; Mismatches 4;
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RESULT 402
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TITLE
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Best Local Similarity
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Best Local
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Local Similarity 93.8%;
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BD201512
Method and reagent
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DNA constructs and methods for screening for increased expression of human apo AI gene
Patent: US 5994061-A 13 30-NOV-1999;
                                                                                                                                                           15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BD201511 17-JUL-2003 Method and reagent for treating diseases or conditions concerning molecule participating in vasculogenic response.
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                                                                                                                                                                                                                                                                                                                                                                                               PC A61P35/00,A61P43/00,C12N5/10,C12N9/00//A61K35/76,C12N15/00,C12N5/00
                                                                                                                                                                                                                                                                                                                                                                                                                                             C12N15/09, A61K31/7088, A61K31/7125, A61K48/00, A61P3/10, A61P17/06, PC
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27-MAR-1998 US 60/079678
PAMELA A PAVCO, ELISABETH ROBERTS, THALE JARVIS, CLAIRE COESHOTT,
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                                                                                                                                                                                                                                                                                                                                   participating in vasculogenic response
Key Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                  concerning molecule
                                                                                                                                                                                                                    /organism="Homo sapiens"
/mol_type="genomic RNA"
/db_xref="taxon:9606"
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Location/Qualifiers
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/mol_type="unassigned DNA"
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Pred. No. 2.7e+02;
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Pred. No. 2.7e+02;
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AUTHORS
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VERSION
KEYWORDS
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BD258338
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ACCESSION
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                                             OS EUKATYOTE
PN JP 2002547195-A/6131
PD 10-DEC-2002
PD 11-APR-2000 JP 2000611654
PR 12-APR-1999 US 60/129390
PI LAWRENCE BLATT, MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN
C12N15/09, A61X38/00, A61X48/00, A61P43/00, A61P43/00, C12N5/10, PC
C12P21/02,
                                                                                                                                                                        1 (bases 1 to 17)
Blatt, L., Zwick, M., Pavco, P. and Mcswiggen, J.
Regulation of repressor genes using nucleic acid molecules
Patent: JP 2002541795-A 6131 10-DEC-2002;
RIBOZYME PHARMACEUTICALS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 17)

Pavco, P.A., Roberts, E., Jarvis, T., Coeshott, C. and Mcswiggen, J.A. Method and reagent for treating diseases or conditions concerning molecule participating in vasculogenic response Patent: JP 2002509721-A 4538 02-APR-2002;
   C12P21/02, C12P21/02//A61K31/711, (C12N5/10, C12R1:91), (C12P21/02, C12R1:91),
                                                                                                                                                                                                                                                                                                                                    Regulation of repressor BD258338
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                                                                                                                                                                                                                                                                                                       JP 2002541795-A/6131.
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24-MAR-1999 JP 2000541291
27-MAR-1998 US 60/079678
PAMELA A PAVCO,ELISABETH ROBERTS,THALE JARVIS,CLAIRE COESHOTT,
JAMES A MCSWIGGEN
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JP 2002509721-A/4538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Key
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/mol_type="genomic RNA"
/db_xref="taxon:9606"
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Pred. No. 2.7e+02;
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acid molecules:
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DEFINITION
ACCESSION
VERSION
KEYWORDS
SOURCE
RESULT 405
CQ616719/c
LOCUS
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AUTHORS
TITLE
JOURNAL
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BD258339
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                                                                                                                Query Match
Best Local Similarity
Matches 15; Conservat
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Best Local Similarity 93.8
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1 (bases 1 to 17)

Blatt,L., Zwick,M., Pavco,P. and Mcswiggen,J.

E Regulation of repressor genes using nucleic acid

NAL Patent: JP 2002541795-A 6132 10-DEC-2002;

PN JP 2002541795-A/6137

PD 10-DEC-2007
                                                                                                                                                                                                          Bource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     source
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CC Regulation of repressor genes using nucleic acid molecules Key Source Location/Qualifiers FT source 1.17

/organism='Eukaryote'.
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PD 10-DEC-2002

PD 11-APR-2000 JP 2000611654

PF 11-APR-1999 US 60/129390

PR 12-APR-1999 US 60/129390

PI LAWRENCE BLATT, MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN

PI LAWRENCE BLATT, MICHAEL ZWICK, PAMELA PAVCO, JAMES MCSWIGGEN

C12N15/09, A61K38/00, A61K48/00, A61P43/00, A61P43/00, C12N5/10, PC
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 CQ616719
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JP 2002541795-A/6132.
                                                                                                                                                                                                                                                                                                                                                               C12P21/02,
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                                                                                CTTTGTTTTGTTTTTT 1262
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                                                                                                                                                                                                                /organism='Eukaryote'
                                                                                                                                                                /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Pred. No. 2.7e+02;
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Sequence 1460 from Patent
CQ616720
CQ616720.1 GI:41666938
                             Homo sapiens (human)
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CQ616719
CQ616719.1 GI:41666937
Eukaryota; Metazoa;
Mammalia; Eutheria;
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Mammalia; Eutheria;
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Primates;
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Sequence 2271 from Patent
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CQ617531.1 GI:41667749
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Shannon,M.E.
                                                                 1 (bases 1 to 17)
Pavco,P., McSwiggenJ., Stinchcomb,D. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 2545 12-FEB-2002;
Location/Qualifiers
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Myosin-like gene expressed in human heart and muscle Patent: WO 0192524-A 2271 06-DEC-2001;
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Mammalia; Eutheria; Primates;
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
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                        /mol_type="unassigned DNA"
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Pred. No. 2.7e+02;
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Beigelman, L., Burgin, A., Beaudry, A., Karpeisky, P.
Matulic-Adamic, J., Sweedler, D. and Zinnen, S.
Synthetic ribonucleic acids with RNAse activity
Patent: US 6528640-A 312 04-MAR-2003;
Location/Qualifiers
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                     AR397930
Sequence 311 from patent
AR397930
                                                                                                                                                                                                                                       Unclassified.

(bases 1 to 17)

(bases 1 to 17)

Pavco,P., McSwiggen,J.A., Stinchcomb,D.T. and Escobedo,J.

Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6566127-A 1069 20-MAY-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unknown.
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/mol_type="unassigned
                                                                                                                                                                                           /mol_type="unassigned RNA"
                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                       organism="unknown"
           GI:40135323
                                                                                                                                                                                                                                                                                                                                              GI:33709475
                                                                                                                                                           1.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.0%; Score 14.4; DB 1; 93.8%; Pred. No. 2.7e+02; tive 0; Mismatches 1;
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AR457783/c
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Sequence 1459 from patent
AR457782
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Unknown.
                                        predominantly in heart and muscle Patent: US 6686188-A 1460 03-FEB-2004; Location/Qualifiers
                                                                          1 (bases 1 to 17)
Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W. Shannon,M.E.
Polynucleotide encoding a human myosin-like polypeptide (
                                                                                                                                 Unknown.
Unclassified.
                                                                                                                                                                                      AR457783
Sequence 1460 from patent
AR457783
AR457783.1 GI:42692840
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Gu,Y., Penn,S.G., Hanzel,D.K., Rank,D.R., Chen,W. Shannon,M.E.
Polynucleotide encoding a human myosin-like polypeptide e predominantly in heart and muscle predominantly in heart and muscle Patent: US 6686188-A 1459 03-FEB-2004;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unknown.
Unclassified
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Beigelman,L., Burgin,A.B., Beaudry,A., Karpeisky,A.,

Matulic-Adamic,J., Sweedler,D. and Zinnen,S.

Oligoribonucleotides with enzymatic activity

Patent: US 6617438-A 311 09-SEP-2003;

Location/Qualifiers
                                                                                                                                                               Unknown.
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/organism="unknown"
/mol_type="genomic
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KEYWORDS
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AX674744
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Best Local Similarity 93.8%;
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AX674744 17 bp DN Sequence 3189 from Patent WO03004526.
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AR458593
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Sequence 2271 from patent
AR458594
                                                                                                                                                                                                                    Polynucleotide encoding a human myosin-like polypeptide expressed predominantly in heart and muscle Patent: US 6686188-A 2271 03-FEB-2004;
                                                                                                                                                                                                                                                                   1 (bases 1 to 17)
Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R.,
                                                                                                                                                                                                                                                                                                          Unknown.
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Patent: US 6686188-A 2270 03-FEB-2004;
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Gu,Y., Ji,Y., Penn,S.G., Hanzel,D.K., Rank,D.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unknown.
                                                                                                                      h 1.0%; Score 14.4; DB 1;
Similarity 93.8%; Pred. No. 2.7e+02;
15; Conservative 0; Mismatches 1
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                                                                                                                                                                                                                                                                                            Unclassified.
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                                                                                                                                                                   /organism="unknown"
/mol_type="genomic DNA"
                                                                                                                                                                                                        Location/Qualifiers
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Pred. No. 2.7e+02;
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Sequence 1717 from Patent
AX730083
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 1189 from Patent
AX729555
AX729555.1 GI:30508898
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses and their use as
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AX674744.1 GI:29333092
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                 Homo sapiens
                                                                                  AX730083.1 GI:30509426
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                 Homo sapiens (human)
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Pred. No. 2.7e+02;
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                           1 (bases 1 to 18)
Brysch, W. and Schlingensiepen, K.
BRYSCH, W. ANTISENSE OLIGONUCLEOTIDE PREPARATION METHOD
PATENTE. WO 9833904-A 1526 06-AUG-1998;
BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
LOCATION/Qualifiers
                                                                                                                                                                                                           Sequence 1526 from Patent A89378
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AX761696
AX761696.1 GI:32256312
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Molecular Engines Laboratories (FR)
Location/Qualifiers
                                                                                                                                             unidentified
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AX662307/c
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AR106885
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Patent: WO 02059293-A 46 01-AUG-2002;
Forster, Anthony C. (US); Blacklow,
                                                                                                                                                                                                                                                  synthetic construct synthetic construct
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Cowsert, L.M., Bennett, C.Frank. and O'Malley, B.W.
Antisense modulation of SRA expression
Patent: US 6107092-A 46 22-AUG-2000;
                                                                                                                                                                                                         Process and compositions for peptide, protein and
                                                                                                                                                                                                                   Forster, A.C. and Blacklow, S.C.
                                                                                                                                                                                                                                       artificial sequences.
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S Schlingensiepen, K.H. and Brysch, W.
An antisense oligonucleotide preparation method
L Patent: JP 2001511000-A 1526 O7-AUG-2001;
BICGNOSTIK GESELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH
OS UDANOWN
P 2001511000-A/1526
PD 07-AUG-2001
PF 30-JAW-1998 JP 1998532533
PF 31-JAW-1998 JP 1998532533
PF 31-JAW-1997 EP 97101531.8
PI KARL HERMANN SCHLINGENSIEPEN, WOLFGANG BRYSCH
PC C12M15/11, CO7H21/04, A61K31/70
CC An antisense oligonucleotide preparation method FH
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1 (bases 1 to 18)
Ohashi, Y., Mitsuhara, I., Sasaki, T., Nagamura, Y., It, H., Iwai, T. and Hiraga, S.
                                                                                                                                                                                                                                                                                       Rice peroxidase having various characteristics
Patent: WO 0142475-A 27 14-GUN-2001;
JAPAN AS REPRESENTED BY DIFFERENCE GENERAL OF MINISTRY OF AGRICULTURE
FORESTRY AND FISHERIES NATIONAL INSTITUTE OF AGROBIOLOGICAL
RESOURCES, KENICHI NOGUCHI YUKO OHASHI,ICHIRO MITSUHARA,TAKUJI
SASAKI,YOSHIAKI NAGAMURA, HIROYUKI ITO, TAKAYOSHI IWAI,SUSUMU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rice peroxidase having various BD096088
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BD066891 18
An antisense oligonucleotide
BD066891 GI:22612494
JP 2001511000-A/1526.
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                                           HIROYUKI ITO,
TAKAYOSHI IWAI,SUSUMU HIRAGA
C12N15/53,C12N9/08,C12Q1/68
                                                                                                                   14-JUN-2001
08-DEC-2000 WO 2000JP008728
10-DEC-1999 JP 99P 352472
YUKO OHASHI,ICHIRO MITSUHARA,TAKUJI SASAKI,YOSHIAKI NAGAMURA,
                                                                                                                                                                                                                  Oryza sativa (rice)
WO 0142475-A/27
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An antisense oligonucleotide preparation method Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Location/Qualifiers
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Pred. No. 2.5e+02;
Location/Qualifiers
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                         artificial sequences.

1 (bases 1 to 20)

S Besterman, J.M., MacLeod, A.R. and Siders, W.M.
Modulation of gene expression by combination the
Patent: JP 2002528391-A 47 03-SEP-2002;
METHYLGENE INC
OS Artificial Sequence
PN JP 2002528391-A/47
PD 03-SEP-2002
PF 19-OCT-1999 JP 2000576885
PF 19-OCT-1999 US 60/104804
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Sequence 5 from Patent W09423026.
A40129
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GENSET (FR)
Other publication AU 6432094 941024
Other publication FR 2703053 940930.
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Vasseur,M., Blumenfeld,M., Meguenni,S. and Poddevin,B.
STAPLE AND SEMI-STAPLE OLIGONUCLECTIDES, METHOD OF PREPARATION AND APPLICATIONS
                                                                                                                                                                                                 synthetic construct
synthetic construct
                                                                                                                                                                                                                               BD244919.1 GI:33054689 JP 2002528391-A/47.
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PF 19-0CT-1999 JP 2000576885
PR 19-0CT-1998 US 60/104804
PI JEFFREY M BESTERMAN, ALAN ROBERT MACLEOD, WILLIAM M SIDERS
A61K48/00, A61K31/165, A61K31/19, A61K31/513, A61K31/517, A61K31/ P
                                                                                                                                                                                                                                                                            Modulation of gene expression
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/organism='Oryza sativa (rice)'.
location/Qualifiers
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/mol_type="genomic DNA"
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Pred. No. 2.5e+02;
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AR298254
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                                                                  l (bases 1 to 20)
Griffals,R., Hoiseth,S.K., Zagursky,R.J.,
Sankaran,B. and Fletcher,L.D.
Chlamydia pneumoniae polynucleotides and u
Patent: US 6559294-A 6476 06-MAY-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                 1 (bases 1 to 20)
Cohen, D., Chumakov, I. and Blumenfeld, M.
Biallelic markers for use in constructing
disequilibrium map of the human genome
Patent: US 6537751-A 9989 25-MAR-2003;
                                                                                                                                                                                        Sequence 6476 from AR315939
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A61X31/7068,A61K31/7088,A61K31/7125,A61K45/00,A61P35/00,C12N15/ PC
                                                                                                                             Unclassified.
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
                   /mol_type="genomic DNA"
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                                                                                                                                                                                                      Boekenkamp,D., Hoppe,H.U. and Burgstaller,P. Detection system for separating constituents production and use of the same Patent: WO 0071747-A 35 30-NOV-2000; Aventis Research & Technologies GmbH & Co. KC
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Allikmets,R., Anderson,K.L., Dean,M., Leppert,M., Lewis,R.A.,
Li,Y., Lupski,J.R., Nathans,J., Rattner,A., Shroyer,N.F., Singh,N.,
Smallwood,P. and Sun,H.
Smallwood,P. and sun acid sequences for ATP-binding cassette
transporter and methods of screening for agents that modify
ATP-binding cassette transporter
patent: US 6713300-A 114 30-MAR-2004;
Location/Qualifiers
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Sequence 114 from patent
AR491020
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milarity 93.8%;
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                                                                                                                   /note="Beschreibung der kunstlichen
Sequenz:Erkennungssystem"
                                                                                                                                            organism="synthetic construct"
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/mol_type="genomic I
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Sequence 6 from Patent WO0071703.
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Sequence 2046 from Patent
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Sequence 15 from Patent WO0071703.
AX053091
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Location/Qualifiers
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                                                    AX184293.1 GI:15135639
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/mol_type="unassigned DNA"
/db xref="taxon:32630"
/noTe="synthetic oligonucleotide"
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/mol_type="unassigned DNA"

/db_xref="taxon:32630"
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Sequence 51
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Sequence 51
AX546302
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Patent: WO 0142511-A 2046 14-JUN-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH
Biotherapeutics Corporation (CA)
                                                                  Besterman, J.M., Macleod, A.R. and Siders, W.M. Modulation of gene expression by combination Patent: EP 1243289-A 51 25-SEP-2002;
                                                                                                                                           synthetic construct synthetic construct
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                      Methylgene, Inc.
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llarity 88.2%;
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/organism="synthetic construct"
/mol_type="unassigned DNA"
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                                                                                                                                                                     1 (bases 1 to 19)
Brysch, W.D. and Schlingensiepen, K.D.
An antisense oligonuclectide preparation method
Patent: EP 0856579-A 562 05-AUG-1998;
                                                                                                                                                                                                                                                                    Sequence 562 from Pat
A90381
A90381.1 GI:6738895
                                                                                                                                                                                                                                                                                                                                                                                                                     16;
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Brysch, W. and Schlingensiepen, K.
ANTISENSE OLIGONUCLEOTIDE PREPARATION METHOD
Patent: WO 9833904-A 562 06-AUG-1998;
BIOGNOSTIK GES (DE); BRYSCH WOLFGANG (DE)
Location/Qualifiers
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l Similarity 93.8%;
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                                                                                                                                              BIOGNOSTIK GES (DE)
Location/Qualifiers
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llarity 84.2%;
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562 from Patent WO9833904.
                                                                                           /organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"
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/note="oligonucleotide"
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/db_xref="taxon:32644"
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                                                           Score 14.2; DB 1;
Pred. No. 2.7e+02;
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Pred. No. 2.7e+02;
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Pred. No. 2.3e+02;
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PD 17-DEC-2002

PP 28-NOV-2001 JP 2001362595

PP 04-MAY-2001 TW 90110785

PR 04-MAY-2001 TW 90
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Sequence
AR294423
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Cohen, D., Chumakov, I. and Blumenfeld, M.
Biallelic markers for use in constructing
disequilibrium map of the human genome
Patent: US 6537751-A 6158 25-MAR-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                      Unknown.
Unknown.
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1 (bases 1 to 19)

Ling,C., Lin,R., Yoo,Z., Huang,X., Lee,B., Lee,S., Lin,Y.,

Huang,C., Heu,H., Shi,C., Yeh,C., Cao,Y. and Pan,C.

Method and detector for identifying subtypes of human papiloma

Patent: JP 2002360271-A 587 17-DEC-2002;

KING CAR FOOD INDUSTRIAL CO LTD
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Oligonuclectide M806108 for identifying HPV CP8061. FH Key
Location/Qualifiers
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17-DEC-2002
28-NOV-2001 JP 2001362595
04-MAY-2001 TW 90110785
CHING-YEE LING, RUEY-WEN LIN, ZHOU-MENG YOO, XIN-HSUAN HUANG, BOW-HAENG LEE,
     Conservative
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic
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Score 14.2; DB 1;
Pred. No. 2.7e+02;
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Pred. No. 2.7e+02
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AX362738/c
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AX362738
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Sequence 99
AX352893
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Ribozyme therapy for the treatment of proliferative
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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                                                                                                                                                                           /note="position 65"
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19 bp
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Pred. No. 2.7e+02;
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Pred. No. 2.7e+02;
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BD065927.1 GI:22611530
JP 2001511000-A/562.
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Sequence 558 from Patent
AX742755
AX742755.1 GI:30576744
1 (bases 1 to 19)
Schlingensiepen, K.H.
                                         unidentified
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King Car Food Industrial Co., Ltd. (T
Location/Qualifiers
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Method and detector for identifying subtypes of human papilloma
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                                                                                                                                                                                                                                                                               /note="Oligonucleotide for Identifying HPV CP8061"
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   and Brysch, W.
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1520 AAAAAAAAAAAAGTAAAAG 1538
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Sequence 57 from Patent WO9813490.
A70736
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Vasseur,M., Blumenfeld,M., Meguenni,S. and Poddevin,B.
STAPLE AND SEMI-STAPLE OLIGONUCLECTIDES, METHOD OF PREPARATION AND APPLICATIONS
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Patent: JP 2001511000-A 562 07-AUG-2001;
BIOGNOSTIK GESELLSCHAFT FUR BIOMOLEKULARE DIAGNOSTIK MBH
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Other publication FR 2703053 940930
Location/Qualifiers
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KARL HERMANN SCHLINGENSIEPEN, WOLFGANG BRYSCH
C12N15/11,C07H21/04,A61K31/70
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30-JAN-1998 JP 1998532533
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/mol_type="unassigned DNA"
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Sequence 57 from Patent EP0834561.
A79220
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 l Similarity
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Dean,N.M., McKay,R., Miraglia,L. and Baker,B.
Antisense oligonucleotide compositions and methods for inibition of C-Jun and C-Fos
Patent: US 598558-A 125 16-NOV-1999;
                                                                                                                                                           Unknown
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/mol_type="unassigned DNA"
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11 Similarity 84.2%;
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Unclassified.
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AR086311
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ARI29739
ARI29739.1 GI:14117636
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                                                                         Sequence 7
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MCKAY,R., Butler,M.M., Wyatt,J. and Cowsert,L.M.

Antisense modulation of pepck-cytosolic expression

Patent: US 6187545-A 143 13-FEB-2001;
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Dean, N.M., McKay, R.,
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/mol_type="unassigned
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                                                                                     from patent US
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Pred. No. 2.5e+02;
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Sequence
AR158717
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1 (bases 1 to 20)
Shannon, K.W., Wolber, P.K., Delenstarr, G.C., Kincaid, R.H.
                                            l Similarity
16; Conserv
                                                                                                                                                                                                                                                          AR158718
Sequence 340 from
AR158718
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                                                                                                                                        Unclassified.
1 (Dases 1 to 20)
Shannon, K.W., Wolber, P.K., Delenstarr, G.C., Kincaid, R.H.
Method for evaluating oligonucleotide probe Patent: US 6251588-A 340 26-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Method for evaluating oligonucleotide probe sequences Patent: US 6251588-A 339 26-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unknown
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              TCTTTGTTTTGTTTTTAA 1264
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/mol_type="unassion«
                                                                                                                           Location/Qualifiers
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/mol_type="unassigned DNA"
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Pred. No. 2.5e+02;
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Sequence 12
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Sequence 6
AR169510
AR169510.1
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AR163731
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Bartel, P.L. and Tavtigian, S.V.
MMSC2--an MWAC1 interacting protein
Patent: US 6291173-A 6 18-SEP-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 20)
Bennett, C. Frank. and Cowsert, L.M.
Antisense inhibition of cytohesin-2
Patent: US 6271029-A 18 07-AUG-2001;
Location/Qualifiers
Unclassified.
1 (Dases 1 to 20)
Dean,N.M., McKay,R., Miraglia,L. and Baker, Antisense oligonucleotide compositions and modulation of activating protein 1
Patent: US 6312900-A 125 06-NOV-2001;
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                     Patent: JP 1993244995-A 7 24

KYOWA HAKKO KOGYO CO LTD

OS Artificial gene

OC Artificial sequence; Gene

OS Zea maize

PN JP 1993244995-A/7

PD 24-SEP-1993

PF KOMATSU YUKI, KIKUCHI

PC C1201/68 C12N15/11;

CC strandedness: Single;

CC topology: Linear;

CC hypothetical: No;

CC anti-sense: No.
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Dean,N.M., McKay,R., Miraglia,L. and Baker,B.
Antisense oligonucleotide compositions and methods for the modulation of activating protein 1
Patent: US 6312900-A 132 06-NOV-2001;
Location/Qualifiers
                                                                                                                                                                                                                                             synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 20)
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JP 1993244995-A/7.
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16; Conservative
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JP 1993244995-A/7
24-SEP-1993
24-SEP-1991 JP 1991243122
KOMATSU YUKI, KIKUCHI YASUHIRO
C1201/68,C12N15/11;
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/mol_type="unassigned DNA"
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E44159/c
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CC Topology: Linear;
FH Key
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FT fource
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                                                                                                                                                                        E44159 20 bp DNA linear PAT 27-AUG-2002 Methods of identification and specific detection of slow-growing mycobacteria by using characteristic base sequence occurring in DNA
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                                                                                                                                                                                                                                                                                                                                 16;
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Hideki,K. and Senshu,U.
Method for analyzing DNA fragment
Patent: JP 1999196874-A 7 27-JUL-1999;
HITACHI LTD
                     mycobacteria by using characteristic base sequence occurring in DNA
                                     artificial sequences.

1 (bases 1 to 20)

Kasai,H., Ezaki,T. and Harayama,S.
Methods of identification and specific detection of slow-growing
                                                                                           synthetic construct
synthetic construct
                                                                                                                        E44159.1 GI:22553300
JP 2001128679-A/2.
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E44159
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gyrase gene
Patent: JP :
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16; Conservative
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Similarity 84.2%;
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C12N15/09, C12Q1/68, G01N27/447, C12N15/00, G01N27/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-JAN-1998 JP 1998005399
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JP 1999196874-A/7
27-JUL-1999
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for analyzing DNA fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism='Unidentified'.
                                                                                                                                                                                                                                                                                                                                                                                /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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2001128679-A 2 15-MAY-2001;
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84.2%; Pred. No. 2.5e+02;
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Pred. No. 2.5e+02;
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PAT 18-JUN-2001
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RESULT 463
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synthetic construct
artificial sequences
1 (bases 1 to 20)
Hirose, K. and Yoshida, T.
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OS Artificial Sequence
PN JP 2001128679-A/2
PD 15-MAY-2001
PF 02-NOV-1999 JP 1999312525
PI HIROAKI KASAI TAKAYUKI EZAKI SHIGEAKI HARAYAMA PC C12N15/09,C12Q1/04,C12Q1/68,C12N15/00
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JP 2000342265-A/9.
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Pred. No. 2.5e+02;
0; Mismatches 3;
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Raben, N., Nichols, R., Plotz, P. and Leff, R.
Assay using recombinant histidyl-tRNA synthetase
Patent: US 5484703-A 7 16-JAN-1996;
Location/Qualifiers
                                                            l Similarity
16; Conserv
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Raben,N., Nichols,R., Plotz,P. and Leff,R.
Raben,N., Nichols,R., Plotz,P. and Leff,R.
Assay using recombinant histidyl-tRNA synthetase
Patent: US 5663066-A 7 02-SEP-1997;
Location/Qualifiers
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Unclassified.
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Sequence 7 from patent US
I63487
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Unclassified.
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Zonana, J., Ferguson, B.M., Headon, D. and Overbeek, P.

Hypohidrotic ectodermal dyplasia genes and proteins

Patent: US 6355782-A 51 12-MAR-2002;

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                      Unknown
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                                                                                                                                                                                                                                                                                                   Unclassified.
                                                                                                                                                                                                                                                                                                                                            Unknown.
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    ACCTGGAGATTGACGCAGA 1036
                                                                Conservative
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/mol_type="unassigned DNA"
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/mol_type="unassigned
                                                                                                                                       /organism="unknown"
/mol_type="unassigned
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84.2%;
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Pred. No. 2.5e+02;
0; Mismatches 3
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                                                            Score 14.2; DB 1;
Pred. No. 2.5e+02;
0; Mismatches 3
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Indels

0

Gape

0

linear

PAT 20-APR-2002

linear

PAT 07-OCT-1997

0

Gaps

0

Indels

0

Gaps

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AR313596/c
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AR224778
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AR307902
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Matches 16; Conserv
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Bennett, C.F. and Freier, S.M.

Antisense modulation of glioma-associated Patent: US 6440739-A 83 27-AUG-2002;

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AR224778
Sequence
AR224778
                                                                                           AR313596
Sequence 4133 from patent
AR313596
AR313596.1 GI:31707022
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                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 113 from patent
AR307902
AR307902.1 GI:31698658
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N
Unclassified.

1 (bases 1 to 20)

Griffais,R., Hoiseth,S.K., Zagursky,R.J., Metcalf,B.J., Sankaran,B. and Fletcher,L.D.

Chlamydia pneumoniae polynucleotides and uses thereof
                                                                                                                                                                                                                                                                                                                    Antisense modulation of raidd expression Patent: US 6551826-A 113 22-APR-2003; Location/Qualifiers
                                                           Unknown
                                                                                                                                                                                                                                                                                                                                                                                            Unknown
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ilarity 84.2%;
Conservative
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                                                                                                                                                                                                                                                                                   /organism="unknown"
/mol_type="genomic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="genomic DNA"
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Pred. No. 2.5e+02;
0; Mismatches 3
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Pred. No. 2.5e+02;
0; Mismatches 3;
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AR371268
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AR314906/c
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Sequence 4 i
AR371268
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                                         l Similarity
16; Conserv
N
                                                                                                                               Unclassified.
1 (Dases 1 to 20)
1 (Dases 1 to 20)
Buchardt,O., Egholm,M., Nielsen,P.E.
Peptide nucleic acids
Patent: US 6395474-A 4 28-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                             16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unclassified.
1 (bases 1 to 20)
Griffais, R., Hoiseth, S.K., Zagursky, R.J.,
Sankaran, B. and Fletcher, L.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AR314906
Sequence 5443 from patent
AR314906
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Location/Qualifiers
                                                                                                                                                                                       Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                     Chlamydia pneumoniae polynucleotides and Patent: US 6559294-A 5443 06-MAY-2003; Location/Qualifiers
                                                                                                                                                                                                   Unknown.
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Similarity 84.2%;
16; Conservative
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                   АЛАЛАЛАЛАЛАСТАЛАЛ 1538
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AAAAGAAAAAAAAAAA 20
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                                                                                 /organism="unknown"
/mol_type="genomic |
                                                                                                                    Location/Qualifiers
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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                                                    1.0%;
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                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                    Score 14.2;
Pred. No. 2.
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Pred. No. 2.
                                           Mismatches
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6395474.
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6559294.
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                                                                                                                                                       Berg, R.H.
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RESULT 471

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REFERENCE
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AUTHORS
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AR491100
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Sequence
AR489489
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Sequence
AR428075
                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 20)
Buchardt,O., Egholm,M., Nielsen,P.E.
Peptide nucleic acid synthons
Patent: US 6710163-A 4 23-MAR-2004;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 20)
Spear, P.G., Warner, M.S., Geraghty, R.J., Martinez, W.M.,
Montgomery, R.I., Cohen, G.H., Eisenberg, R.J., Whitbeck, C.J.
Krummenacher, C.
Cellular proteins which mediate herpesvirus entry
Cellular proteins which mediate herpesvirus entry
Patent: Us 6641818-A 5 04-NOV-2003;
Location/Qualifiers
 1 (bases 1 to 20)
Buchardt, O., Buchardt, D., Egholm, M., Nielsen, P.E.
Synthetic procedures for peptide nucleic acids
Patent: US 6713602-A 4 30-MAR-2004;
Location/Qualifiers
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                                                                                                                 Sequence 4 from patent US AR491100
                                                                                                                                                                                                                                                                                                                                                                                        Unknown
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                                                                                                                                                                                                                                                                                      /organism="unknown"
/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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                                                                                                       GI:47258960
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Pred. No. 2.5e+02
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Sequence 15 from Patent WO0105435.
AX078001 GI:13157746
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                                                                                            /mol_type="unassigned
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="genomic
                                                                                                                 organism="Homo sapiens"
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Antisense therapy for hormone-regulated tumors
Patent: WO 0105435-A 15 25-JAN-2001;
THE UNIVERSITY OF BRITISH COLUMBIA (CA) ; Miyake, Hideaki (JP)
Location/Qualifiers
                                                                                                                                                    Antisense therapy for hormone-regulated tumors Patent: WO 0105435-A 15 25-JAN-2001; THE UNIVERSITY OF BRITISH COLUMBIA (CA) ; Miyal Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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RESULT 478
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Matches 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                   synthetic construct
synthetic construct
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Identification method and specific detection method
mycobacteria utilizing dna gyrase gene
Patent: EP 1098003-A 3 09-MAY-2001;
MARINE BIOTECHNOLOGY INSTITUTE CO., LTD. (JP)
Location/Qualifiers
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Sequence 3 from Patent |
AX137428
AX137428.1 GI:14273633
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ilarity 84.2%;
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/mol type="unassigned DNA"
/mol type="azon:32630"
/noTe="Synthetic DNA"
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|db_xref="taxon:32630"
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AX462672/c
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AX452909/c
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Bernard, M., Sourdille, P. and Guyomarch, H. Microsatellite markers from Triticum tauschii Patent: EP 1217079-A 416 26-JUN-2002;
                                                                                                                                          Sequence 416 from Patent AX462672
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                                                                                                                              AX462672.1
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ORGANISM
Aegilops tauschii
Aegilops tauschii
Bukaryota; Viridiplantae; Streptophyta; Em
Spermatophyta; Magnoliophyta; Liliopsida;
Pooideae; Triticeae; Aegilops.
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Mammalian nuclear receptor cofactors cf7
Patent: WO 0242322-A 10 30-MAY-2002;
LION Bioscience AG (DE)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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/db_xref="taxon:32630"
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Pred. No. 2.5e+02;
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                                Embryophyta; Tracheophyta;
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AX462737/c
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Eukaryota; Viridiplantae; Streptophyta; Embryophyt
Spermatophyta; Magnoliophyta; Liliopsida; Poales;
Pooideae; Triticeae; Aegilops.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 481 from AX462737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
16; Conserv
                                                                                                        Full-length cDNA sequences
Patent: EP 1308459-A 3983 07-MAY-2003;
                                                                                                                                              Isogai,T., Sugiyama,T., Otsuki,T., Wakamatsu,A., Sato,H., Ishii,S.,
Yamamoto,J.I., Isono,Y., Hio,Y., Otsuka,K., Nagai,K., Irie,R.,
Tamechika,I., Seki,N., Yoshikawa,T., Otsuka,M., Nagahari,K. and
                                                                                                                                                                                                                                                         AX750458
Sequence 3983 from Patent
AX750458
AX750458.1 GI:32132876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bernard, M., Sourdille, P. and Guyomarch, H. Microsatellite markers from Triticum tauschii Patent: EP 1217079-A 481 26-JUN-2002; INSTITUT NATIONAL DE LA RECHERCHE AGRONOMIQUE
                                                                                                                                                                                                                   synthetic construct
                                                                                                                                                                                                                                 synthetic construct
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                                                                                                                                                                                                       artificial sequences.
                                                                                             Helix Research Institute (JP)
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           /organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Aegilops tauschii"
/mol_type="unassigned DNA"
/db_xref="taxon:37682"
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/db_xref="taxon:37682"
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note="an artificially synthesized primer sequence"
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Pred. No. 2.5e+02;
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Susceptibility locus for schizophrenia
Susceptibility locus for schizophrenia
Patent: WO 03050301-A 98 19-JUN-2003;
Gurling, Hugh Malcolm Douglas (GB)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
AX785487
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                                                                                                                                                                                    recognition Patent: WO 03060160-A 1139 24-JUL-2003;
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Similarity 84.2%;
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Similarity 84.2%;
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                                                                   Similarity
                             TGTTGCTGCTGCCTTTGTG 747
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Oreochromis niloticus (Nile tilapia)
Oreochromis niloticus
Oreochromis niloticus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Teleostei; Euteleostei; Neoteleostei;
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes;
                                                                                                                                                                                                                                                 Lie,Y., Slettan,A., Hoeyum,M. and Lingaas,F. Verification of food origin based on nucleic
                                                                            /organism="Oreochromis niloticus"
/mol_type="unassigned DNA"
/db_xref="taxon:8128"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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                 Bynthetic construct

artificial sequences.

1 (bases 1 to 20)

RS Takagi.M., Shinji.H. and Ota,K.

Peptide having a function regulating transcription of gene

Patent: JP 2001269176-A 302-OCT-2001;

AL Patent: JP 2001269176-A/3

PD 2001269176-A/3

PD 20-OCT-2001

SCHOOL BECOME

PN JP 2001269176-A/3

PD 02-OCT-2001

PF 27-MAR-2000 JP 200087536

PF 47-MAR-2000 JP 200087536

PF 27-MAR-2000 JP 200087536

PC C12N15/09,C07K14/415,C12N1/15,C12N1/19,C12N1/21,C12N5/10// PC

C12N15/09,C12R1:91),(C12N5/10,C12R1:91),C12N15/00,C12N5/00,

PC (C12N15/00,C12R1:91),(C12N5/10,C12R1:91)

CC Description of Artificial Sequence: Synthetic primer DNA FH

Location/Qualifiers.
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         Location/Qualifiers.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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                                                                                     TITLE
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synthetic construct

artificial sequences.

E 1 (bases 1 to 20)

E 1 (bases 1 to 20)

S Takagi, M., Shinji, H. and Ota, K.

Peptide having a function regulating transcription of ger

Patent: JP 2001269176-A 5 02-OCT-2001;

AGENCY OF IND SCIENCE & TECHNOL

OS Artificial Sequence

PN JP 2001269176-A/5

PD 02-OCT-2001
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PC (C12N19
PC (C12N19
CC Descrip
Key
                                                                                                               synthetic construct artificial sequences.
1 (bases 1 to 20)
                                                      A method of arraying genome clone Patent: JP 2001321190-A 1783 20-NOV-2001; THE INSTITUTE OF PHYSICAL AND CHEMICAL RE
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Similarity 84.2%;
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Description of Artificial Sequence: Synthetic primer DNA FH Location/Qualifiers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-MAR-2000 JP 2000087536
MASARU TAKAGI,HIDEAKI SHINJI,KEN OTA
C12N15/09,C07K14/415,C12N1/15,C12N1/19,C12N1/21,C12N5/10// PC
Artificial Sequence
JP 2001321190-A/1783
20-NOV-2001
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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Pred. No. 2.5e+02;
0; Mismatches 3
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Pred. No. 2.5e+02;
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                                                           AND CHEMICAL RESEARCH, YUGENKAISHA
                                                                                                                                                                                                                     clone
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RESULT 490
S47176S4/c
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AUTHORS
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VERSION
KEYWORDS
SOURCE
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BD161924
                DEFINITION
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C12N15/00,
C12N15/
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                                                                                                                                                  1 Similarity
16; Conserv
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PD 09-JUL-2002
PF 27-DEC-2000 JP 2000399573
PF 27-DEC-2000 JP 2000399573
PI MICHIFUMI TANGA, HIROSHI OKAMURA, KOJIRO TAKAHASHI PC C12N15/09,C12N15/09 CC Mei C12N15/09,C12N15/09 CC Mei C12N15/09 out thermal cycle of PCR using DNA- CC
                                                                                                                                                                                                                                                                                                                                                                             Patent: JP 2002191369-A 1 09-JUL-2002;
TOYO KOHAN CO LTD,KOJIRO TAKAHASHI
OS Artificial Sequence
PN JP 2002191369-A/1
PD 09-JUL-2002
PF 27-DEC-2000 JP 2000399573
S47176S4 20 by lipoprotein lipase (introns 3, segment 4 of 5).
                                                                                              N
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1 (bases 1 to 20)
Tanga,M., Okamura,H. and Takahashi,K.
Method for carrying out thermal cycle
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BD161924.1 GI:27867682
JP 2002191369-A/1.
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Method for carrying out thermal
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EIICHI SOEDA
CI2N15/09,C12N15/09,C12M1/00,C12Q1/68,G01N33/53,G01N33/566,
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Pred. No. 2.5e+02;
0; Mismatches 3;
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Pred. No. 2.5e+02;
0; Mismatches 3;
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             NA linear PRI 08-MAY-1993
8} [human, Genomic, 20 nt,
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AUTHORS
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AB068086/c
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SOURCE
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TITLE
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S47179.1
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Synthetic construct DNA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GenBank staff at the National Library of Medicine created this entry [NCBI gibbsq 117241] from the original journal article. Regions surrounding three polymorphic sites.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 20)

Gotoda, T., Yamada, N., Murase, T., Shimano, H., Shimada, M., Harada, K., Kawamura, M., Kozaki, K. and Yazaki, Y.

Detection of three separate DNA polymorphisms in the human lipoprotein lipage gene by gene amplification and restriction
                                                                                                                                                                                                                                                                                                                chromosome 1p35-p36
Genomics 74 (1), 55-70
21269192
                                                                                                                                                                                                                                                                                                                                                                                                     Chen,Y.Z., Hayashi,Y., Wu,J.G., Takaoka,E., Maekawa,K.,
Watanabe,N., Inazawa,J., Hosoda,F., Arai,Y., Mizushima,H.,
Morohashi,A., Ohira,M., Nakagawara,A., Liu,S., Hoshi,M., Horii,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                 Submitted (04-AUG-2001) Akira Horii, Tohoku University School of Medicine, Molecular Pathology; 2-1 Seiryomachi, Aoba-ku, Sendai, Miyagi 980-8575, Japan (E-mail:horii@mail.cc.tohoku.ac.jp, Tel:81-22-717-8042, Fax:81-22-717-8047)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        synthetic
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J. Lipid Res. 33 (7), 1067-1072 (1992)
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                                                                                                                                                                                                                                           Direct Submission
                                                                                                                                                                                                                                                             Horii, A.
                                                                                                                                                                                                                                                                                                                                                                      A BAC-based STS-content map spanning a 35-Mb region of human
                                                                                                                                                                                                                                                                                                                                                                                         and Soeda, E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 artificial sequences.
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/note="forward primer for human STS sts-H96854 at 1p36 sts-H96854 obtained from clones B293A18, B122E3, B91D18, Human BAC library RPCI-11"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                     /db_xref="taxon:32630"
                                                                                          /organism="synthetic construct"
/mol_type="genomic DNA"
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Pred. No. 2.5e
0; Mismatches
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